# A multi-scale model reveals cellular and physiological mechanisms underlying hyperpolarisation-gated synaptic plasticity

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### 13 Abstract

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- <sup>14</sup> Neurons in the medial vestibular nucleus (MVN) display hyperpolarisation-gated synaptic plasticity,
- <sup>15</sup> where inhibition believed to come from cerebellar cortical Purkinje cells can induce long-term
- <sup>16</sup> potentiation (LTP) or long-term depression (LTD) of vestibular nerve afferent synapses. This
- phenomenon is thought to underlie the plasticity of the vestibulo-ocular reflex (VOR). The
- 18 molecular and cellular mechanisms involved are largely unknown. Here we present a novel
- <sup>19</sup> multi-scale computational model, which captures both electrophysiological and biochemical
- <sup>20</sup> signalling at vestibular nerve synapses on proximal dendrites of the MVN neuron. We show that
- 21 AMPA receptor phosphorylation at the vestibular synapse depends in complex ways on dendritic
- 22 calcium influx, which is in turn shaped by patterns of post-synaptic hyperpolarisation and
- vestibular nerve stimulation. Hyperpolarisation-gated synaptic plasticity critically depends on the
- activation of LVA calcium channels and on the interplay between CaMKII and PP2B in dendrites of
- <sup>25</sup> the post-synaptic MVN cell. The extent and direction of synaptic plasticity depend on the strength
- <sup>26</sup> and duration of hyperpolarisation, and on the relative timing of hyperpolarisation and vestibular
- <sup>27</sup> nerve stimulation. The multi-scale model thus enables us to explore in detail the interactions
- <sup>28</sup> between electrophysiological activation and post-synaptic biochemical reaction systems. More
- <sup>29</sup> generally, this model has the potential to address a wide range of questions about neural signal
- <sup>30</sup> integration, post-synaptic biochemical reaction systems and plasticity.

31

## 32 Introduction

- <sup>33</sup> Inhibition in neural circuits plays a fundamental role in modulating the activity and dynamic
- responsiveness of neurons (Brock et al., 1952; Isaacson and Scanziani, 2011; Gidon and Segev,
- <sup>35</sup> 2012; Bloss et al., 2016; Doron et al., 2017; Hull, 2017). In cerebellum-dependent motor learning,
- <sup>36</sup> inhibitory projections of Purkinje neurons in the cerebellar cortex to neurons in the vestibular
- <sup>37</sup> nucleus (VN) and deep cerebellar nucleus (DCN), are essential for the induction of adaptive plasticity
- <sup>38</sup> in extra-cerebellar motor pathways. Purkinje neurons integrate afferent signals from a wide range
- <sup>39</sup> of sensory systems and inputs from inferior olive neurons, which encode 'errors' in ongoing motor

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Present address: <sup>‡</sup>Tri-Institutional <sup>8</sup> Training Program in Computational <sub>9</sub> Biology and Medicine, New York, USA <sup>10</sup> acts (*Ito, 2002; Yakusheva et al., 2007; Wulff et al., 2009*). While initial models proposed that motor

<sup>41</sup> memories were acquired and retained within the cerebellar cortex (*Ito, 1982*), much experimental

and theoretical evidence now shows that motor learning involves the transfer of cortically-acquired
 motor memories to sub-cortical and extra-cerebellar structures for consolidation and retention

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47 particular. The VOR is the reflex by which the eyes move in the opposite direction to the head 48 in order to maintain a stable image. The VOR is a robust and tractable experimental system in

<sup>49</sup> which to study cerebellum-dependent motor learning (*Blazquez et al., 2004*: *Boyden et al., 2004*:

<sup>50</sup> du Lac et al., 1995: Miller et al., 2005: Broussard et al., 2011: Shin et al., 2014). Errors in the gain

of the VOR, where the evoked eve movements are larger or smaller than those required to precisely

s2 compensate for head movements, result in a slippage of the optical image on the retina and the

<sup>53</sup> loss of visual field stability. In response to 'retinal slip' errors encoded by inferior olive neurons,

54 Purkinje cells in the cerebellar flocculus are believed to induce potentiation or depression of the

vestibular nerve afferent synapses on medial VN (MVN) neurons, so increasing or decreasing the

<sup>56</sup> gain of the VOR appropriately to cancel retinal slip (*Raymond and Lisberger, 1998; Medina, 2010*;

57 Clopath et al., 2014; Carcaud et al., 2017).

Recent studies have revealed a distinct form of hyperpolarisation-gated synaptic plasticity in MVN and DCN neurons, where inhibition presumed to be mediated by Purkinje cell synapses, induces long-term potentiation (LTP) or depression (LTD) of heterologous excitatory, glutamatergic synapses on these neurons (*Pugh and Raman, 2009; McElvain et al., 2010*). This represents therefore a plausible cellular mechanism by which cerebellar Purkinje cells may induce lasting changes in sub-cortical neural networks during the consolidation of a learned motor memory (*Kassardjian et al., 2005; Shutoh et al., 2006; Anzai et al., 2010*).

The cellular and molecular mechanisms that mediate hyperpolarisation-gated plasticity are 65 largely unknown. In both DCN and MVN neurons, excitatory synaptic stimulation paired with a pat-66 tern of hyperpolarisation and release from hyperpolarisation, has been shown to induce LTP in the 67 excitatory synapses (Pugh and Raman, 2009: McElvain et al., 2010). In DCN neurons the activation 68 of low-threshold T type calcium channels (LVCa channels) upon the release of hyperpolarisation, is 69 required for LTP (*Person and Raman, 2010*). T type calcium channels are also expressed in MVN 70 neurons (Serafin et al., 1991a.b: Him and Dutia, 2001: Engbers et al., 2013), and interestingly these 71 channels are significantly up-regulated during vestibular compensation, the behavioural recovery 72 that takes place after vestibular deafferentation (Him and Dutig, 2001; Straka et al., 2005; Menzies 73 et al., 2010). 74

Vestibular nerve and Purkinie cell synapses have recently been demonstrated to be closely 75 apposed on dendrites of parvocellular MVN neurons, providing the anatomical substrate for a 76 close spatial interaction between convergent inhibitory and excitatory synapses in these neurons 77 (Matsuno et al., 2016). To investigate the cellular and molecular mechanisms that might underlie 78 such interactions, we developed a novel multi-scale model. This model integrates the electrophysio-79 logical model of a Type B MVN neuron developed by *Ouadroni and Knopfel (1994*) with biochemical 80 models of postsynaptic calcium signalling and the subsequent activation of LTP- or LTD-inducing 81 pathways, originally designed to model plasticity in hippocampal dendritic spines (Stefan et al., 82 2008: Li et al., 2012: Mattioni and Le Novère, 2013). The resulting multi-scale model allowed us to 83 explore the molecular mechanisms invoked by the patterns of inhibition and excitatory stimulation 84 that mediate hyperpolarisation-gated synaptic plasticity. We show that different types of synaptic 85 activity are associated with different sources of calcium influx, with low-voltage activated calcium 86 channels (LVACCs) becoming the dominant route of calcium entry during hyperpolarisation-gated 87 synaptic plasticity. We also show that AMPA receptor phosphorylation (which we use as a readout 88 for synaptic potentiation) shows a highly sigmoidal response to calcium, and that this response is

mainly determined by the balance between kinase and phosphatase activity. We further show that

- <sup>91</sup> the extent and direction of hyperpolarisation-gated synaptic plasticity depend on the strength and
- <sup>92</sup> duration of the hyperpolarising stimulus, as well as on the relative timing between hyperpolarisation
- <sup>93</sup> and excitation. Our multi-scale model provides a novel in-silico test bed for further investigations
- <sup>94</sup> into the interplay between inhibition, excitation and biochemical processes in synaptic plasticity.
- 95 Results

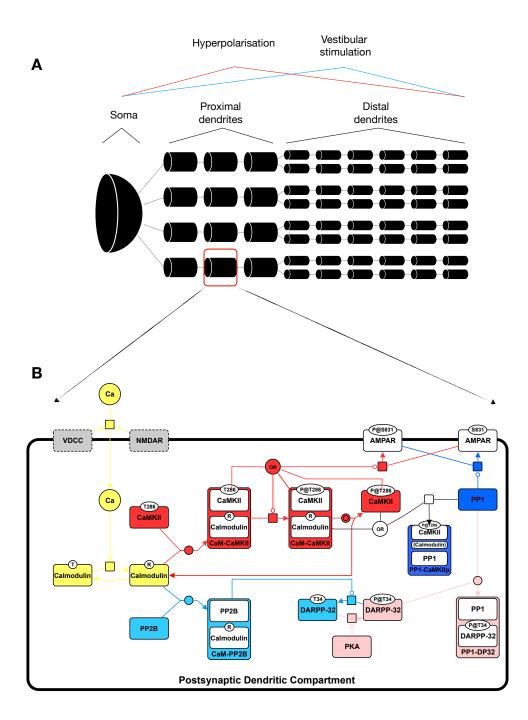
## 96 Biochemical Pathway Model of Synaptic Plasticity

We modelled the biochemical pathways underlying synaptic plasticity based on an earlier model 97 by Li et al. (2012) (see Methods for details). A Systems Biology Graphical Notation (SBGN) diagram 98 (Le Novère et al., 2009) describing the main components of our model is shown in Fig. 1B. Calcium enters through voltage-dependent calcium channels (VDCCs) and NMDA receptors (shown in grey, 100 not explicitly included in the chemical model). Calcium then binds to and activates calmodulin 101 (shown in yellow). Active (R-state) calmodulin can then activate either the calmodulin-dependent 102 protein kinase II (CaMKII) pathway (shown in red) or the Protein phosphatase 2B (PP2B) (also known 103 as calcineurin (CaN)) pathway (shown in blue). CaMKII is active when it is bound to calmodulin or 104 autophosphorylated. Active CaMKII phosphorylates AMPA receptors, leading to enhanced AMPAR 105 activity. In the PP2B pathway, calmodulin-bound PP2B dephosphorylates DARPP-32, and thereby 106 releases PP1 from inhibition. Active PP1 can dephosphorylate AMPA receptors, counteracting the 107 effect of CaMKII. PP1 also directly dephosphorylates CaMKII itself. The PKA pathway (shown in 108 pink) can reduce dephosphorylation of AMPA receptors by activating DARPP-32. Taken together. 100 the model captures the sequence of events leading from calcium influx to changes in AMPAR 110 phosphorylation state. 111

# AMPA receptor phosphorylation shows a complex dependency on calcium concen tration

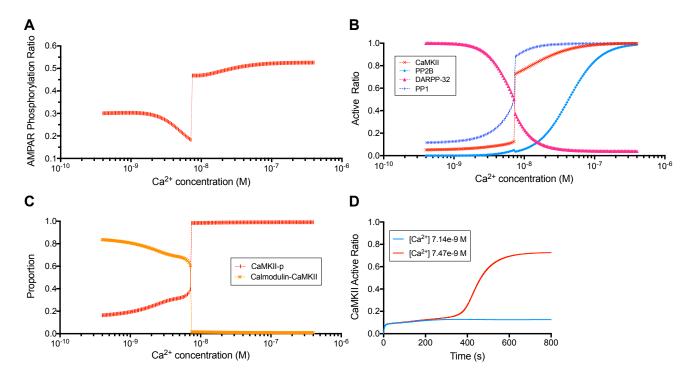
<sup>114</sup> In order to establish the usefulness of our biochemical model as a model of synaptic plasticity, we <sup>115</sup> first examined the dependence of AMPA receptor phosphorylation on calcium concentration. To <sup>116</sup> this end, we determined the effects of a step change in  $[Ca^{2+}]$  from baseline at equilibrium to a <sup>117</sup> range of concentrations over the range  $4 \times 10^{-10}$  M to  $4 \times 10^{-7}$  M (Fig. 2). The time course of changes <sup>118</sup> in the active molecular species in the system were observed over 1600s after the step change in <sup>119</sup>  $[Ca^{2+}]$ . As output, we computed the AMPAR phosphorylation ratio (phosphorylated AMPAR over <sup>120</sup> total AMPAR).

As can be seen from Fig. 2A, the relationship between [Ca<sup>2+</sup>] and AMPAR phosphorylation is 121 complex. At very low calcium concentrations there is a baseline level of AMPAR phosphorylation 122 of about 30%. Moderate increases in calcium concentrations (to the nanomolar range) lead to a 123 decrease in AMPAR phosphorylation ratio. At  $[Ca^{2+}]$  of between 7 and 7.5  $\times$  10<sup>-9</sup> M however, there is 124 a sharp increase in AMPAR phosphorylation ratio, so that at  $[Ca^{2+}]$  values of 7.5  $\times$  10<sup>-9</sup> M and higher 125 about 50% of total AMPA receptors are phosphorylated at the end of the 1600s simulation period. 126 In order to identify the molecular determinants of the changes in AMPA receptor phosphoryla-127 tion, we examined the activation ratios (active over total concentrations) of other key molecular 128 species in the system over the same range of  $Ca^{2+}$  concentrations (Fig. 2B). The activation patterns 120 of kinases and phosphatases show sigmoidal responses to initial calcium concentration, with dif-130 ferent directions, incline, and transition concentrations. At sub-threshold Ca<sup>2+</sup> concentrations, the 131 dominant active form of CaMKII is calmodulin-bound (non-phosphorylated) CaMKII, whereas at 132 higher Ca<sup>2+</sup> phosphorylation becomes the main contributor to CaMKII activity (Fig. 2C). In order to 133 highlight the bi-stable response pattern of CaMKII, we ran time courses at calcium concentrations 134 just below and just above the threshold concentration at which the system switches (Fig. 2D). 135 The bi-stable response is also present when calmodulin concentration is reduced, although the 136 sharpness of the response and the calcium concentration at which the system changes between 137 states varv (Fig. S1). 138



**Figure 1.** Simplified schematic of the models used in this work. **A**. NEURON model with 61 electrical compartments, representing the soma, proximal dendrites, and distal dendrites. **B**. Chemical model with the key molecular species and reactions that underlie calcium-dependent synaptic plasticity.

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**Figure 2.** Dependence of AMPAR phosphorylation and relative activation ratios of key molecular species on  $[Ca^{2+}]$ . **A**. AMPAR phosphorylation ratio (phosphorylated AMPAR over total AMPAR) as a function of  $[Ca^{2+}]$  over the range  $4 \times 10^{-10}$  M to  $4 \times 10^{-7}$  M. For each data point the biochemical model was re-initialised and after reaching equilibrium the intracellular  $[Ca^{2+}]$  was stepped up to successively higher values (logarithmic scan of  $[Ca^{2+}]$  range, 151 simulations). Each simulation was run for 1600 s, to steady state. Note the initial dephosphorylation of AMPA receptors with increasing  $[Ca^{2+}]$ , which is followed by a switch to a large sustained increase in AMPAR phosphorylation when  $[Ca^{2+}]$  is increased above a threshold value for inducing long-term potentiation. **B**. Relative activation ratios (concentration of active molecules divided by total molecules) for CaMKII, PP2B, DARPP-32, and PP1 as a function of  $[Ca^{2+}]$ . **C**. The proportion of the two active CaMKII forms: CaMKII-p (phosphorylated CaMKII) and calmodulin-bound (non-phosphorylated) CaMKII as a function of  $[Ca^{2+}]$ . **D**. Time-course of activation of CaMKII over a period of 800 s, at two values of  $[Ca^{2+}]$  on either side of the threshold concentration.

Since dephosphorylation of AMPA receptors is associated with LTD and AMPA receptor phospho-139 rylation with LTP, this behaviour of the biochemical system is consistent with the widely accepted 140 model by which a moderate increase in  $[Ca^{2+}]$  will result in the depression of excitatory synapses. 14 and a greater increase will result in their potentiation (Lisman, 1989; Malenka, 1994). We therefore 142 used AMPA receptor phosphorylation ratio as a readout in our subsequent simulation experiments. 143 We next implemented the biochemical model within dendritic compartments of the MVN neuron. 144 in order to study LTD and LTP of the excitatory vestibular nerve synapses that impinge upon the 145 MVN neuron. 146

# Dendritic [Ca<sup>2+</sup>] profiles induced by vestibular synaptic stimulation with and with out hyperpolarisation

We next studied how different stimulation protocols affect calcium dynamics in the proximal dendrites of the MVN neuron. For this, we used a multi-compartmental electrophysiological model of an MVN Type B neuron (*Quadroni and Knopfel, 1994*; *Graham et al., 2009*) in the NEURON software (*Carnevale and Hines, 2006*) (Fig 1A). We stimulated each neuronal compartment in the model by either activating the excitatory vestibular synapses (Vestibular Stimulation protocol, "VS") or by activating the excitatory vestibular synapses together with hyperpolarisation of the cell membrane (Hyperpolarisation + Vestibular Stimulation protocol, "H+VS").

The results are shown in Fig. 3. The MVN neuron had a resting firing rate of around 24 Hz before 156 vestibular stimulation, similar to the resting activity observed in experimental studies (Iohnston 157 et al., 1994: McElvain et al., 2010) (Fig. 3 A). Stimulation of the vestibular nerve input at 100 Hz 158 ("VS protocol") evoked a firing rate of around 56 Hz, similar to the response evoked by vestibular 159 nerve stimulation in the study of *McElvain et al.* (2010). There was a significant increase in [Ca<sup>2+</sup>] 160 for the duration of the vestibular nerve stimulation, with the result that the total dendritic  $[Ca^{2+}]$ 161 was elevated to 2.3 times the baseline level (Fig. 3 B, D). Analysis of the individual dendritic ion 162 channel currents (Fig. 3C) showed that at rest, the  $Ca^{2+}$  influx was mediated predominantly by 163 LVA  $Ca^{2+}$  channels and by HVA  $Ca^{2+}$  channels that were activated during each action potential. By 164 contrast during VS stimulation, the activation of synaptic NMDA channels in addition to the LVA and 165 HVA channels, led to a significant increase of dendritic  $Ca^{2+}$  influx for the duration of the vestibular 166 stimulation (Fig. 3C. D). 167

As shown in Fig. 3B, in response to the H+VS stimulation protocol there was an initial hyperpo-168 larisation of around 25 mV over the first 250 ms, which was followed by a rebound depolarisation 169 and increase in firing rate upon the release of inhibition. Dendritic  $[Ca^{2+}]$  decreased initially during 170 the period of inhibitory stimulation, but the large influx of  $[Ca^{2+1}]$  during the rebound depolarisation 171 resulted in the net dendritic  $[Ca^{2+}]$  being elevated to a level 3.64 times the baseline (Fig. 3D). The 172  $Ca^{2+}$  influx from LVA channels activated upon the release of inhibition was the predominant source 173 of dendritic Ca<sup>2+</sup> during the H+VS stimulation. Together with the NMDA receptor mediated influx. 174 this led to the highly elevated levels of dendritic  $[Ca^{2+1}]$  (Fig. 3D). 175

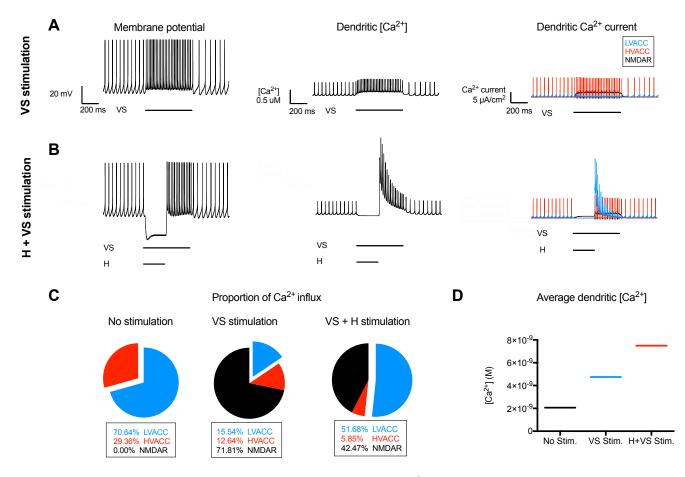
Together, these results demonstrate that during vestibular nerve stimulation alone (VS protocol) the activation of synaptic NMDA receptors causes a moderate elevation of dendritic  $[Ca^{2+}]$ , while pairing hyperpolarisation with vestibular nerve stimulation in the H+VS protocol led to the elevation of dendritic  $[Ca^{2+}]$  to a substantially higher level. Thus, postsynaptic hyperpolarisation has a marked effect on dendritic  $[Ca^{2+}]$  influx, when the activation of LVA channels coincides with activation of the synaptic NMDA receptors.

## Multi-scale modelling reveals molecular pathways activated in response to vestibu lar synaptic stimulation with and without hyperpolarisation

<sup>184</sup> We next examined the effect of calcium influx under the VS and H+VS protocols on downstream <sup>185</sup> signalling pathways. In order to do this, we combined the NEURON and COPASI models described

<sup>185</sup> signalling pathways. In order to do this, we combined the NEURON and COPASI models described
 <sup>186</sup> above into one multi-scale modelling pipeline. Specifically, we first used the NEURON model to

simulate the VS and H+VS protocols, as described above. The readout from the neuron model



**Figure 3.** Effects of VS and H+VS stimulus protocols on membrane potential and  $[Ca^{2+}]$  in a dendritic compartment of a Type B MVN cell. The membrane potential and action potential firing (left panel), total  $[Ca^{2+}]$  profiles (middle panel) and  $Ca^{2+}$  currents dynamics (right panel) in a single dendritic compartment of a NEURON model of a Type B MVN cell in response to vestibular nerve synapse stimulation at 100 Hz for 550 ms (VS protocol, **A**), and to vestibular stimulation combined with membrane hyperpolarisation for the first 250 ms of the stimulation period (H+VS protocol, **B**). **C**. the contribution of low-voltage activated calcium channels (LVACC), high-voltage activated calcium channels (HVACC) and NMDA receptor channels (NMDAR) to the changes in  $[Ca^{2+}]$  observed in the two stimulus protocols. Note the activation of HVACC during action potential firing, and the activation of synaptic NMDAR during VS stimulation. During H+VS stimulation, the release of the membrane hyperpolarisation results in a large activation of LVACC Ca<sup>2+</sup> currents. **D**. Total dendritic  $[Ca^{2+}]$  averaged over the simulation period for the VS and H+VS stimulus protocols after scaling due to Ca<sup>2+</sup> buffering effect (Fig. 11).

was a profile of intracellular calcium concentration over time. These profiles were then used as 188 an input for a COPASI simulation of the calcium-dependent biochemical signalling pathways as 189 described earlier. The interface between NEURON and COPASI was coded in Python and included 190 two steps to facilitate a smooth conversion from electrical to biochemical model. First, we binned 191 time steps from the NEURON model into larger time intervals, because biochemical simulations 192 happen at a larger time scale and therefore do not require the same temporal resolution as 193 electrophysiological models. Second, we applied a scaling factor in order to account for the fact that 194 a portion of the calcium entering the cell will immediately bind to intracellular calcium buffers not 195 explicitly represented in our biochemical model, and the effective calcium concentration available 196 for calmodulin binding will therefore be smaller than the overall calcium concentration. 197

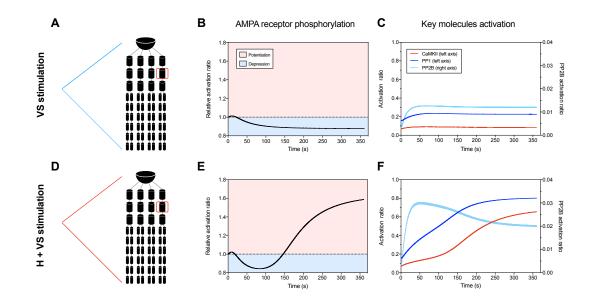
The outcomes of the multi-scale simulations are shown in Fig. 4. The dendritic  $[Ca^{2+}]$  profiles 198 induced by the VS and H+VS stimulation protocols had substantially different effects on the key 190 components of the biochemical model and the resultant phosphorylation of AMPA receptors. In 200 particular the moderate rise of dendritic  $[Ca^{2+}]$  induced by the VS stimulation led to a modest 201 activation of CaMKII and PP2B (Fig. 4 A), and a sustained decrease in the level of AMPA receptor 202 phosphorylation over the course of the simulation (Fig. 4C). By contrast, the greater rise in  $[Ca^{2+}]$ 203 induced by the H+VS stimulation led to a marked activation of PP2B and CaMKII (Fig. 4B). Following 204 an initial decrease in active AMPA receptor phosphorylation ratio, the H+VS stimulation protocol 205 induced a large sustained increase in AMPA receptor phosphorylation level (Fig. 4D). Presumably 206 the rise in  $[Ca^{2+1}]$  evoked by the H+VS protocol was greater than that required to trigger auto-207 phosphorvlation of CaMKII and the consequent high level of phosphorylation of the AMPA receptors, 208 while in the VS protocol the modest rise in  $[Ca^{2+1}]$  was insufficient to trigger this "switch-like" 209 behaviour. The VS and H+VS protocols thus had opposite effects on synaptic AMPA receptors. This 210 is consistent with the experimental observation of synaptic LTD and LTP in the study of *McElvain* 211 et al. (2010). 212

Taken together, vestibular stimulation following hyperpolarisation results in markedly increased calcium influx, activation of calcium-dependent kinases over phosphatases and a resulting net increase in AMPA receptor phosphorylation. In contrast, vestibular stimulation alone results in a more moderate calcium influx, a different balance of kinase and phosphatase activity, and a net dephosphorylation of AMPA receptors.

# LVA channel density determines dendritic calcium dynamics and hyperpolarisation gated synaptic plasticity

To examine further the role of MVN LVA channels in mediating hyperpolarisation-gated plasticity, we 220 explored the relationship between LVA channel density, the post-hyperpolarisation rebound calcium 221 influx and effects on synaptic AMPAR phosphorylation. We compared the effects of the VS and H+VS 222 protocols on AMPAR phosphorylation ratio in a "Type Bhigh IVA neuron" model, in which the dendritic 223 LVA channel density is significantly higher than the normal Type B neuron (Quadroni and Knopfel, 224 1994) (Table 2), and in the hypothetical "Type B<sub>low LVA</sub> neuron" in which the LVA channel density was 225 reduced to 50 % of that in the normal Type B neuron (Table 2). As expected, in the Type B<sub>high IVA</sub> 226 neuron the post-hyperpolarisation rebound firing and the dendritic Ca<sup>2+</sup> influx were significantly 227 greater than in the normal Type B neuron, reflecting the higher density of dendritic LVA channels in 228 this cell type. By contrast the Type Biow 1VA neuron showed a much smaller rebound depolarisation 220 and reduced dendritic  $Ca^{2+}$  influx than the normal Type B neuron (Fig. 5). As shown in Fig. 6, the 230 effects of the VS and H+VS protocols on AMPAR phosphorylation levels in these neuron types were 231 also markedly different. In the Type B<sub>high LVA</sub> neuron, the VS stimulation protocol caused a slightly 232 more pronounced decline in AMPA receptor phosphorylation level than in the "normal" Type B 233 neuron. The H+VS stimulation protocol evoked a larger, more rapid increase in AMPA receptor 234 phosphorylation (Fig. 6). By contrast in the Type B<sub>low LVA</sub> neuron VS stimulation caused a slower, 235 less marked decrease in AMPAR phosphorylation levels than in the normal Type B neuron, while 236 the H+VS stimulation did not result in increased AMPAR phosphorylation, instead producing a slight 237

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**Figure 4.** Time-course of activation of key molecular species and AMPAR phosphorylation in response to VS and H+VS stimulation. Left part is the changes in active ratios of CaMKII, PP1 and PP2B in response to the VS stimulation protocol (**A**) and H+VS stimulation (**B**). Right part corresponding changes in AMPAR phosphorylation in response to the VS stimulation protocol (**A**) and H+VS stimulation protocol (**B**). Note that the VS protocol evokes a moderate activation of the key molecular species and results in a net de-phosphorylation of AMPAR over the period of stimulation, while the H+VS protocol induces a marked activation of PP1 and CaMKII, and results in a sustained phosphorylation of AMPA receptors.

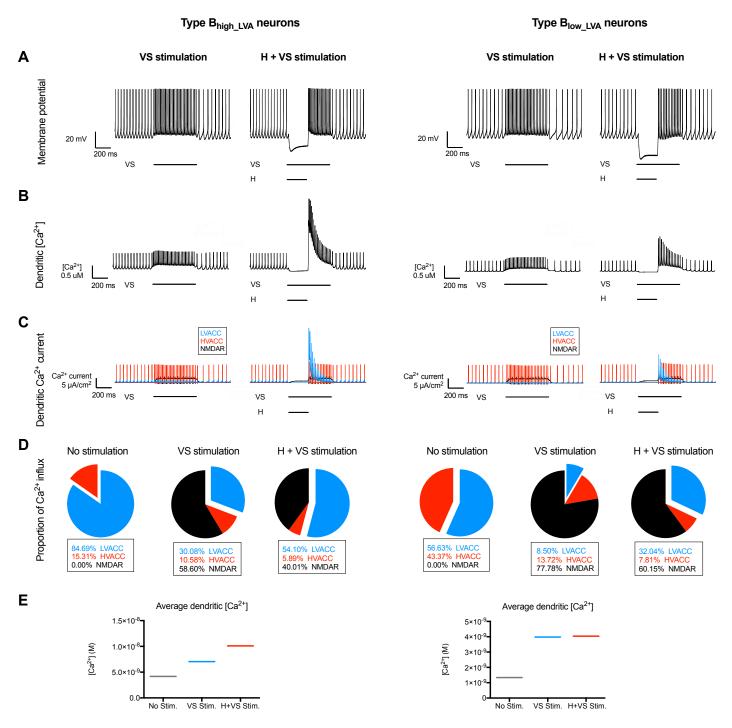
238 decline.

Together, these results indicate that the density of LVA channels is a major determinant of dendritic calcium dynamics and hyperpolarisation-gated synaptic plasticity.

## Hyperpolarisation-gated synaptic plasticity depends on the balance of kinases and phosphatases

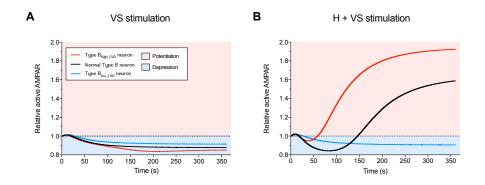
Given the importance of dendritic calcium influx and its biochemical effects in regulating AMPA 243 receptor phosphorylation levels, we next examined the effects of varying the relative concentrations 244 of the two key molecules CaMKII and PP2B, on the response of the normal MVN Type B neuron 24 to the H+VS stimulation protocol (Fig. 7). We ran a total of 9 simulations where we reduced the 246 intracellular concentrations of CaMKII and PP2B to 66% and 33% of the base model independently 247 and in combination, to explore the effects of changing the absolute levels of these molecules as well 248 as their relative ratios. As shown in Fig. 7, total CaMKII concentration is the main determinant of 249 AMPAR phosphorylation level, although the balance between kinase and phosphatase concentration 250 also plays a role. At normally high CaMKII concentrations (Fig. 7A), there is always an increase in 251 AMPAR phosphorylation, though that increase is more pronounced when PP2B concentrations is 252 reduced. Interestingly, when CaMKII concentration is slightly reduced (Fig. 7B), this can produce 253 either an increase or a decrease in AMPAR phosphorylation, depending on PP2B concentration. At 254 low CaMKII levels (Fig. 7C), the kinase activity is no longer enough to induce a net increase in AMAPR 255 phosphorylation at normal or 66 % of normal levels of PP2B. Only if PP2B concentration is also 256 lowered to 33 % of its base model concentration can we see an increase in AMPAR phosphorylation. 257 but this increase is slow and moderate in scale. 258 These findings demonstrate the importance of CaMKII as a key molecular species in mediating 259

hyperpolarisation-gated plasticity, and show that the net effectiveness of the H+VS stimulation
 protocol is critically dependent on the balance between the kinase and its antagonist phosphatase
 PP2B.

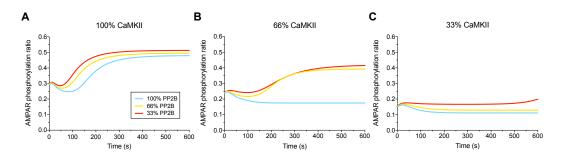


**Figure 5.** Effects of VS and H+VS stimulus protocols on membrane potential and  $[Ca^{2+}]$  in a dendritic compartment of a Type  $B_{high_LVA}$  or Type  $B_{low_LVA}$  MVN cell. The same stimulation protocols and analysis as in Fig. 3 are applied here. In the Type  $B_{high_LVA}$  neuron the post-hyperpolarisation rebound firing and the dendritic  $Ca^{2+}$  influx were significantly greater than in the normal Type B neuron, while the Type  $B_{low_LVA}$  neuron showed a much smaller rebound depolarisation and reduced dendritic  $Ca^{2+}$  influx than the normal Type B neuron. Note that because the types of neurons differ in LVA density, they have different basal calcium concentrations, and a direct comparison of absolute values between types of neurons is therefore difficult.

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**Figure 6.** Effects of VS and H+VS stimulation protocols on AMPAR phosphorylation depend on LVA channel expression in MVN neurons. Effects of the VS (**A**) and H+VS (**B**) stimulation protocols on AMPAR phosphorylation in a normal Type B MVN neuron, a Type B neuron expressing a high level of LVACC (Type B <sub>high\_LVA</sub> cell), and a hypothetical class of MVN neuron with low expression of LVACC (Type B <sub>low LVA</sub> cell).



**Figure 7.** Dependence of AMPAR phosphorylation response on CaMKII, PP1 and PP2B levels. Effects of varying the concentrations of CaMKII and PP2B in a normal Type B MVN neuron on the phosphorylation of AMPAR in response to the H+VS stimulation protocol. Simulations were run where CaMKII and PP2B concentrations were set to 100 %, 66 % and 33 % of those typically found in a neuron.

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## Hyperpolarisation-gated plasticity depends on the strength, duration, and relative timing of the hyperpolarising stimulus

Finally we explored what features of the hyperpolarising stimulus are important for determining the strength and direction of synaptic plasticity. We investigated this by simulating repeated combinations of inhibitory stimuli for 600 s each. Across simulations, we varied either the strength, duration, or relative timing of the hyperpolarising stimulus (corresponding to 600, 600 and 400 inhibition/excitation pairs in 600 s), while keeping the other two parameters and the parameters governing the excitatory stimulus constant.

We first varied the amplitude of the hyperpolarising stimulus. In order to do this, we used our multi-scale model to simulate a H+VS protocol in normal Type B MVN neurons with hyperpolarisation strengths ranging from 0 pA to 600 pA (Fig. 8A). In all cases, each excitatory pulse lasted for 550 ms. Each hyperpolarising stimulus started at the same time as the excitatory stimulus and lasted for 250 ms.

As shown in Fig. 8B, both dendritic  $Ca^{2+}$  concentration and AMPA receptor phosphorylation 276 displayed a complex dependency on the strength of the hyperpolarising stimulus. Small amounts 277 of membrane hyperpolarisation caused essentially no change (or a small degrease) in calcium 278 levels, and no change in AMPA receptor phosphorylation. Intermediate hyperpolarisation strengths 270 resulted in a moderate increase in [Ca<sup>2+</sup>] and a decrease in AMPA receptor phosphorylation. Once 280 hyperpolarisation strength reaches a threshold of 450 pA, the system switches: CaMKII reaches 281 sustained levels of activation, resulting in an overall increase of AMPA receptor phosphorylation. 282 (Fig. 8D). 283

Next, we varied the duration of the hyperpolarising stimulus. In order to do this, we used our
 multi-scale model to simulate a H+VS protocol in normal Type B MVN neurons with hyperpolarisa tion duration ranging from 0 ms to 550 ms (Fig. 9A). In all cases, we used a hyperpolarising stimulus
 of 475 pA, which started at the same time as a 550 ms excitatory stimulus.

The dendritic [Ca<sup>2+</sup>] response and AMPA receptor phosphorylation showed a complex depen-288 dence on the duration of the H stimulus (Fig. 9B). A short period of hyperpolarisation (less than 289 80 ms) leads to a small increase of  $[Ca^{2+}]$  in the rebound and no change or a decrease in AMPA 290 receptor phosphorylation. With very long periods of hyperpolarisation that overlap with the excita-291 tory stimulation, the net Ca concentration can even be lower than that without stimulation. This will 292 also lead to a decrease in AMPAR activation. At hyperpolarisation durations in between, there is 293 a robust rise in calcium levels, and a stable "up" state is reached at which almost half of all AMPA 29/ receptors are phosphorylated (Fig. 9D). 295

Finally we varied the timing between hyperpolarising and depolarising stimuli. In order to do this, we used our multi-scale model to simulate a H+VS protocol in normal Type B MVN neurons. We varied the onset of the hyperpolarising stimulus from -500 ms to +250 ms from the onset of the excitatory stimulus (Fig. 10A). In all cases, the amplitude of the hyperpolarising stimulus was 475 pA, and the duration was 250 ms.

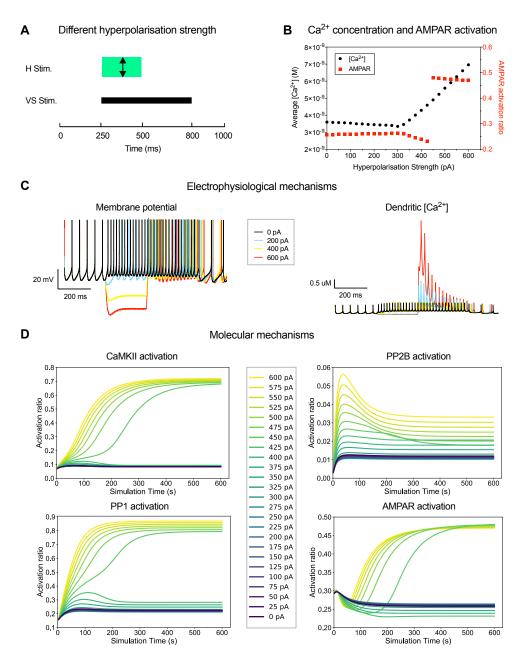
The dendritic [Ca<sup>2+</sup>] response showed a marked dependence on the relative timing of the hyperpolarising and excitatory stimuli (Fig. 10B). A maximal elevation of [Ca<sup>2+</sup>] was seen when the release of the membrane hyperpolarisation corresponded with the start of the VS stimulation. In contrast, dendritic [Ca<sup>2+</sup>] was at its lowest when the onsets of hyperpolarising and excitatory stimuli coincided. Correspondingly, there is a narrow window of relative timings where strong and sustained AMPA receptor phosphorylation is observed (Fig. 10B, D), suggesting that the precise timing of hyperpolarisation and excitation is crucial for hyperpolarisation-gated synaptic plasticity.

### 308 Discussion

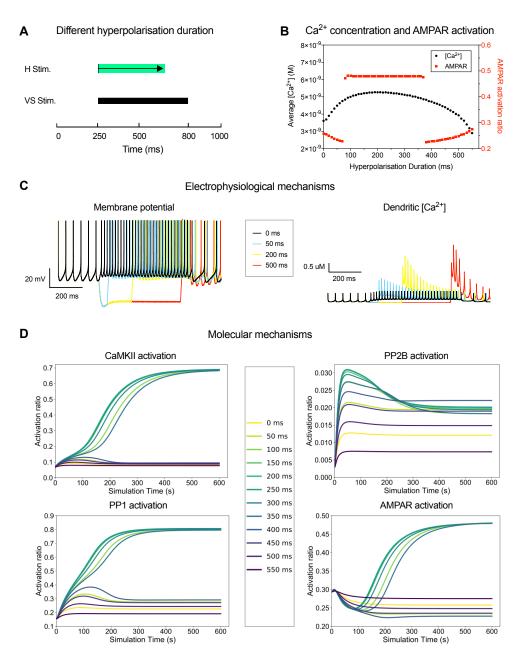
<sup>309</sup> In this study we developed a detailed multi-scale model incorporating electrophysiological and

<sup>310</sup> biochemical processes that regulate AMPA-receptor phosphorylation in MVN neurons, to investigate

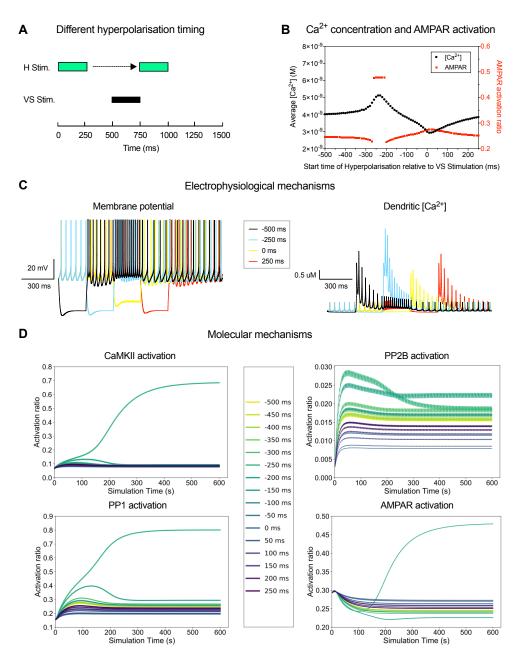
the interactions between membrane hyperpolarisation and synaptic plasticity. The electrophysio-



**Figure 8.** Effects of varying hyperpolarisation strength on AMPA receptor activation. **A**. Simulation set-up: Hyperpolarising stimuli of various strengths were applied to MVN type B neurons. **B**. AMPAR activation at the end of each 600 s simulation plotted as a function of average  $[Ca^{2+}]$  during the same simulation. **C**. Electrophysiological mechanisms (dynamics of the membrane potential and dendritic  $[Ca^{2+}]$ ) for selected hyperpolarisation strengths. **D**. Molecular changes (activation of kinases and phosphatases and AMPAR phosphorylation over time) for a range of hyperpolarisation strengths.



**Figure 9.** Effects of varying hyperpolarisation duration on AMPA receptor activation. **A**. Simulation set-up: Hyperpolarising stimuli of varying duration were applied to MVN type B neurons. **B**. AMPAR activation at the end of each 600 s simulation plotted as a function of average  $[Ca^{2+}]$  during the same simulation. **C**. Electrophysiological mechanisms (dynamics of the membrane potential and dendritic  $[Ca^{2+}]$ ) for selected hyperpolarisation durations. **D**. Molecular changes (activation of kinases and phosphatases and AMPAR phosphorylation over time) for a range of hyperpolarisation durations.



**Figure 10.** Effects of varying the timing between hyperpolarisation and excitation on AMPA receptor activation. **A.** Simulation set-up: Hyperpolarising and excitatory stimuli were applied at varying relative timing to MVN type B neurons. **B.** AMPAR activation at the end of each 600 s simulation plotted as a function of average  $[Ca^{2+}]$  during the same simulation. **C.** Electrophysiological mechanisms (dynamics of the membrane potential and dendritic  $[Ca^{2+}]$ ) for selected hyperpolarisation onset timings. **D.** Molecular changes (activation of kinases and phosphatases and AMPAR phosphorylation over time) for a range of hyperpolarisation onset timings.

logical component of the multi-scale model was based on well-established NEURON models which 312 faithfully reproduce the membrane characteristics of rodent Type B MVN neurons in vitro (Ougdroni 313 and Knopfel, 1994). The biochemical component of the multi-scale model was based on models of 314 calcium signalling and the activation of protein kinases and phosphatases in postsynaptic neuronal 315 compartments (Stefan et al., 2008; Li et al., 2012; Mattioni and Le Novère, 2013). Dendritic Ca<sup>2+</sup> 316 concentration profiles obtained from the NEURON model of the MVN neuron were routed to the 317 biochemical component of the model. This allowed us to model changes in the phosphorylation ra-318 tio of AMPA receptors, as a proxy of long-term potentiation or depression of the activated synapses 319 in a dendritic compartment of the MVN neuron. 320

The multi-scale model thus allowed us to examine for the first time the cellular and molecular 321 mechanisms involved in hyperpolarisation-gated synaptic plasticity, where the synaptic strength 322 of vestibular nerve afferent inputs in MVN neurons is regulated by inhibitory inputs presumably 323 from Purkinie cell synapses (*McElvain et al., 2010*). Similar interactions are also seen in DCN 324 neurons (Pugh and Raman, 2009; Person and Raman, 2010), suggesting that hyperpolarisation-325 gated synaptic plasticity is a fundamental mechanism in cerebellum-dependent motor learning 326 We used our model to specifically determine the effects on calcium signalling and AMPA receptor 327 regulation induced by experimental protocols that cause either a potentiation or a depression of the 328 vestibular synapses in MVN neurons (*McElvain et al., 2010*). During resting activity, dendritic Ca<sup>2+</sup> 329 concentration showed small transients largely due to Ca<sup>2+</sup> influx with each action potential through 330 high-threshold  $Ca^{2+}$  channels, leading to low baseline levels of AMPA receptor phosphorylation. A 331 train of synaptic stimulation at 100 Hz (VS stimulation protocol) has previously been shown to induce 332 LTD in vestibular synapses (McElvain et al., 2010). In contrast, if the same train was accompanied 333 by a period of hyperpolarisation (H+VS protocol), then this resulted in LTP (*McElvain et al.*, 2010) 334 Our model reproduced both effects, with VS stimulation alone causing only moderate increases in 335 calcium concentration and a net decrease in AMPA receptor phosphorylation. In contrast, H+VS 336 stimulation evoked larger elevations in calcium concentration, primarily due to the activation of 33 low-voltage activated  $Ca^{2+}$  channels upon the release of hyperpolarisation. This led to a substantial 338 increase in AMPA receptor phosphorylation. The role of low-voltage activated  $Ca^{2+}$  channels in 339 mediating the high dendritic  $[Ca^{2+}]$  following periods of hyperpolarisation is in agreement with the 340 finding of *Person and Raman* (2010) that the induction of hyperpolarisation-gated synaptic plasticity 34 in DCN neurons is dependent on LVCa channel activation. 342

This hyperpolarisation-gated synaptic plasticity relies on the delicate balance between kinase 343 and phosphatase activity in the post-synaptic dendrite. CaMKII and PP2B compete for activation by 344 calcium-activated calmodulin, and their relevant concentrations are instrumental in determining 345 whether LTP or LTD is induced. This is consistent with previous results (Stefan et al., 2008: Li et al., 346 2012). Further downstream, CaMKII and PP1 compete to determine the phosphorylation status 347 both of AMPA receptors and of CaMKII itself. Their bi-stable behaviour determines the bi-stable 348 response of AMPA receptor phosphorylation (Zhabotinsky, 2000: Miller et al., 2005: Pi and Lisman, 349 2008)). This means that the biochemical reaction system functions as a switch that can produce 350 long-term potentiation or long-term depression in response to changes in calcium concentrations. 351 Our present findings show how the electrophysiological activation of the MVN neuron by patterns 352 of synaptic excitation coupled with inhibition, induces changes in dendritic calcium concentrations 353 and drives the biochemical reaction system to bring about hyperpolarisation-gated plasticity at the 354 vestibular nerve synapse. 355

The amount and direction of hyperpolarisation-gated synaptic plasticity are dependent in complex ways on the strength and duration of the hyperpolarising stimulus, as well as the relative timing between hyperpolarisation (inhibition) and excitation. Long-term potentiation is observed when hyperpolarisation is sufficiently strong, and when the release from hyperpolarisation coincides with an excitatory stimulus.

Our multi-scale simulation workflow uses the open source tools NEURON (*Carnevale and Hines*, and COPASI (*Hoops et al., 2006*) that work with community-driven standards (NeuroML

(Gleeson et al., 2010) and SBML (Hucka et al., 2003)). This provides a useful template for the 363 development of further multi-scale models linking electrophysiological and biochemical models 364 more generally. The modularity of our model means it can also be extended to include more 365 components of the biochemical signalling pathways or a wider neuronal signalling network. It 366 thus has the potential to address a wide range of questions about neural signal integration, post-367 synaptic biochemical reaction systems and plasticity. While our current model appears to contain 368 the essential components required to express hyperpolarisation-gated synaptic plasticity, further 369 developments are necessary to incorporate additional inputs to MVN neurons that modulate 370 synaptic plasticity, and cellular mechanisms involved in the consolidation of synaptic plasticity 371 through the regulation of gene expression, for example, 372

## 373 Methods

## <sup>374</sup> Biochemical model of signalling pathways underlying synaptic plasticity

The biochemical model used here is based on previous models of calcium signalling and the 375 activation of protein kinases and phosphatases in postsynaptic compartments (Stefan et al., 2008; 376 Li et al., 2012; Mattioni and Le Novère, 2013), starting from Ca<sup>2+</sup> input and leading to AMPA receptor 377 phosphorylation as a readout. Specifically, we have built on an earlier model by Li et. al (Li et al., 378 2012), which we helped encode in SBMI format, and which is now available on BioModels Database 379 (Li et al., 2010) (BIOMD000000628). The model follows the basic SBGN reaction scheme introduced 380 in Fig 1B. Accounting for the fact that several of the model components can exist in different 381 functional states (Stefan et al., 2014) and modelling each of those states explicitly, there are a total 382 of 129 molecular species and 678 reactions in the model. Initial concentrations of chemical species 383 were taken from the model by *Li et al.* (2012). Concentration of CaMKII was changed to  $1 \times 10^{-5}$  M 384 for MVN neurons (*Biber et al., 1984*). Besides, we added the reaction: CamR Ca2 AC + PP2B  $\rightarrow$ 385 CamR Ca2 AC PP2B, which is necessary for completeness, but was missing in the model by Li 386 et al. (2012). We also added PP2B activation to the model which was described in the paper by 387 Li et al. (2012), but was not included in their supplemental model file. For autophosphorylation 388 of CaMKII, we used the polynomial formula used by Li et al. (2012) to compute phosphorylation 389 rates, which accounts for the fact that autophosphorylation proceeds from one active subunit to its 390 neighbour in the CaMKII holoenzyme. However, we slightly modified the rate formula to ensure 391 that the autophosphorylation rate was always greater than 0, in order to be more biochemically 392 accurate. At last, we removed  $Ca^{2+}$  buffer proteins from the system, because we already account 393 for calcium buffering when translating between electrical and chemical models (see below). 394 Initial concentrations of chemical species were taken from previous literature based on rat 395

<sup>396</sup> brains (*Biber et al., 1984; Li et al., 2012; Mattioni and Le Novère, 2013*) and are presented in table

<sup>397</sup> 1. Reaction rates and other parameters were as described by *Li et al.* (2012).

Name	Concentration (M)	
Calmodulin	$1.00 \times 10^{-4}$	
CaMKII	$7.00 \times 10^{-5}$	
PP2Bi	$6.00 \times 10^{-6}$	
DARPP-32	$3.00 \times 10^{-6}$	
РКА	$1.20 \times 10^{-8}$	
PP1a	$2.00\times10^{-6}$	
AMPAR_p	$1.66 \times 10^{-7}$	
AMPAR	$1.49 \times 10^{-6}$	

**Table 1.** Initial concentration of molecular species in the biochemical model

The biochemical model was edited and run in COPASI 4.16 (*Hoops et al., 2006*). It is available in the project's GitHub repository (https://github.com/YubinXie/multiscale-synaptic-model) in SBML level 2.4 (.xml) format.

## 401 Simulation of Chemical model

In order to observe concentration dynamics of key chemical species over time, the "Time Course"
 function in COPASI was used. We chose a time interval of 0.0001 s, and the "deterministic (LSODA)"
 method. The concentration of Ca<sup>2+</sup> was either fixed throughout the time course, or altered using
 the "Events" functionality in COPASI in order to simulate a dynamic calcium signal (see below).

In order to explore equilibrium behaviours at different  $Ca^{2+}$  concentrations (Fig. 2), we used the "Parameter Scan" function in COPASI. Initial  $Ca^{2+}$  concentrations ranging from  $4 \times 10^{-10}$  M to  $4 \times 10^{-7}$  M were scanned, with 151 logarithmic intervals. At each initial Ca concentration, the simulation started from the initial state of the biochemical model and was run for 1600 s with the time interval being 0.0001 s. This was long enough for the system to reach an equilibrium state. The

value of Ca<sup>2+</sup> concentration was fixed at initial value during the whole simulation.

## 412 Electrical model of MVN type B neuron

We used a multi-compartmental electrophysiological model of an MVN Type B neuron adapted from 413 Ougdroni and Knopfel (1994) and implemented (Graham et al., 2009) in NEURON (Carnevale and 414 Hines, 2006). The model neuron consisted of a soma, 4 proximal dendrites and 8 distal dendrites. 415 comprising 61 electrical compartments. Each compartment included up to nine active ionic channels 416 (Table 2), and with the exception of the soma also included one excitatory vestibular nerve synapse 417 containing by AMPA receptors and NMDA receptors. The strength of the vestibular nerve synapses 418 was adjusted so that stimulation at a frequency of 100 Hz caused an increase in firing rate of the 419 model MVN neuron similar to that observed experimentally by *McElvain et al.* (2010). 420 We used the two canonical subtypes of Type B MVN neurons modelled by Quadroni and Knopfel 421 (1994) in our simulations. The normal Type B MVN neuron, representing the majority of MVN 422 neurons (Straka et al., 2005), expresses a moderate level of low-voltage activated Ca<sup>2+</sup> channels 423 (LVA channels, Table 2). By contrast the "Type B<sub>high LVA</sub>" neuron, normally representing some 10 % of 424 MVN neurons (Serafin et al., 1991a; Him and Dutia, 2001), expresses a higher level of LVA channels 425 and shows a pronounced low-threshold rebound firing ("low-threshold Ca<sup>2+</sup> spike") upon release 426

from hyperpolarisation. While the models of *Quadroni and Knopfel* (**1994**) accurately replicate the electrophysiological properties of normal Type B and Type B<sub>high LVA</sub> neurons, it should be noted that

the expression of LVA  $Ca^{2+}$  channels in MVN neurons and the incidence of post-hyperpolarisation

rebound firing is heterogeneous (*Serafin et al., 1991a; Straka et al., 2005*). Indeed LVA channel

expression in MVN neurons is rapidly upregulated after deafferentation, with the number of Type
 B<sub>high LVA</sub> neurons increasing significantly during vestibular compensation (*Him and Dutia, 2001*;

**Straka et al., 2005**). In this light, to investigate the functional role of LVA channels in the present study, we also modelled a third, hypothetical "Type  $B_{low LVA}$ " MVN neuron, where the LVA Ca<sup>2+</sup>

 $_{434}$  study, we also modelled a time, hypothetical hype  $D_{low_LVA}$  with reducin, where the LV.  $_{435}$  channel density was reduced to 50 % of that in normal Type B neurons (Table 2).

## 436 Multi-scale interface and stimulation protocols

We applied the stimulation protocols that have been shown experimentally to induce bidirectional 437 plasticity of excitatory synapses in DCN and MVN neurons in vitro (Person and Raman, 2010; 438 *McElvain et al., 2010*), to determine the evoked dendritic [Ca<sup>2+</sup>] profiles and their effects on synaptic 439 AMPA receptors in the multi-scale model. The vestibular synapses on the proximal and distal 440 dendritic compartments of the MVN model were activated for 550 ms at a frequency of 100 Hz 441 Repeated periods of vestibular synaptic activation alone ("VS" protocol) have been shown to 447 cause LTD of excitatory synapses in MVN neurons (*McElvain et al., 2010*). Alternatively, in the 443 hyperpolarisation + vestibular synaptic activation protocol ("H+VS"), vestibular synaptic stimulation 444 was paired with hyperpolarisation of the post-synaptic cell for 250 ms, by the injection of inhibitory 449

	Maximal conductance per membrane area ( $\mu Scm^{-2}$ )		
Channels	Soma	Proximal Dendrites	Distal Dendrites
Na	43000	2880	0
Nap	23.6	38	0
Н	66	66	0
bK(fast)	37530	2572	640
AHP	2716	0	0
K(slow)	519	406	0
А	755	0	0
Ca(HVA)	2385	1417	350
Ca(LVA)	166	325.5/651/1607*	50
Na(leak)	37.8	0.7	0.71
Ca(leak)	74.6	1	1
K(leak)	166	3.69	3.68

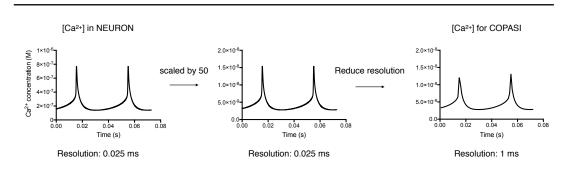
Table 2. Distribution and density of ionic conductances of MVN B neurons(Quadroni and Knopfel, 1994)

\*The values are for Type B<sub>low LVA</sub>, normal Type B, Type B<sub>high LVA</sub> MVN neurons, respectively.

current into all of the dendritic compartments. This pattern of stimulation was shown to induce LTP
 of vestibular nerve synapses in MVN neurons (*McElvain et al., 2010*).

Values of [Ca<sup>2+</sup>] in the mid-proximal dendrites during the normal resting activity of the MVN neuron and in response to the VS and H+VS stimulation protocols were obtained from NEURON, and used as the input to the biochemical model in COPASI. The [Ca<sup>2+</sup>] values obtained from NEURON were scaled down by a factor of 50, to account for Ca buffering and sequestration processes not explicitly included in our biochemical model (Fig.11).

<sup>453</sup> This scaling factor brought the  $[Ca^{2+}]$  values calculated in NEURON into the physiological range of <sup>454</sup> intracellular Ca<sup>2+</sup> concentration known from experimental studies (*Sharma et al., 1995*). In addition, <sup>455</sup> the chemical model has a much lower time resolution than the electrical model, with a simulation <sup>456</sup> time step of 1 ms vs. 0.025 ms in the electrical model. We therefore calculated the average of 40 <sup>457</sup> successive values of  $[Ca^{2+}]$  in the NEURON model, to provide the COPASI model with an input value <sup>458</sup> of  $[Ca^{2+}]$  at every 1 ms time step (Fig.11). After  $[Ca^{2+}]$  scaling and time resolution transformation, <sup>459</sup> the resulting  $[Ca^{2+}]$  values was added to the COPASI input file as an "Event". Then, the biochemical



**Figure 11.** Demonstration of  $[Ca^{2+}]$  conversion from electrical modelling tool NEURON to biochemistry modelling tool COPASI. Firstly, the  $[Ca^{2+}]$  values obtained from NEURON were scaled down by a factor of 50, to account for  $Ca^{2+}$  buffering and sequestration processes not explicitly included in our biochemical model. Then, the time resolution was scaled up by a factor of 40, given the relatively slower dynamics in biochemistry system when compared to the electrical system.

460 model could be run in COPASI. All the process was automatically done by a Python script (provided

<sup>461</sup> here: https://github.com/YubinXie/multiscale-synaptic-model). One thing to note is that, before

adding the stimulation events into COPASI model, the COPASI model should run for 1000 s with

the  $Ca^{2+}$  being the average  $Ca^{2+}$  of the corresponding (Type B, Type  $B_{low_LVA}$ , Type  $B_{high_LVA}$ ) resting neurons.

**Regular H + VS simulation protocol** (Fig. 3, 5, 4, 6): In a 1000 ms window, 100Hz 550 ms VS stimuli starts at 250 ms and ends at 800 ms. 475 pA inhibitory stimuli (hyperpolarisation) starts at 250 ms and ends at 500 ms. This 1000 ms simulation is constantly repeated until the simulation ends (360 s). Inhibitory stimuli of 475 pA on Type  $B_{high_LVA}$  neurons gave a huge [Ca<sup>2+</sup>] rebound, which is too huge to be visualized in a reader-friendly way. In Fig. 5, 6, 320 pA inhibitory stimuli was

used in Type B<sub>high LVA</sub> neurons.

Hyperpolarisation strength simulation protocol (Fig. 8): In a 1000 ms window, a 100 Hz
550 ms VS stimulus starts at 250 ms and ends at 800 ms. Inhibitory stimulus (hyperpolarisation)
with strength from 0 pA to 600 pA starts at 250 ms and ends at 500 ms. This 1000 ms window is
constantly repeated until the simulation ends (600 s).

Hyperpolarisation duration simulation protocol (Fig. 9): In a 1000 ms window, 100 Hz, a
550 ms VS stimulus starts at 250 ms and ends at 800 ms. A 475 pA inhibitory stimulus (hyperpolarisation) with duration from 0 ms to 550 ms starts at the same time. This 1000 ms window is
constantly repeated until the simulation ends (600 s).

**Hyperpolarisation timing simulation protocol** (Fig. 10): To allow a large range of timing scanning, here we used a 1500 ms time window. In this window, a 100 Hz 250 ms VS stimulus starts

at 250 ms and ends at 500 ms. A 250 ms inhibitory stimulus (hyperpolarisation) with strength 475 pA

starts ranging form 500 ms earlier to 250 ms later than the VS stimulus. This 1500 ms simulation is
 constantly repeated until the simulation ends (600 s).

<sup>484</sup> Individual simulations were run on standard desktop and laptop computers. Sets of simulations <sup>485</sup> for figures 8 to 10 were run on the Eddie Compute Cluster at the University of Edinburgh.

## 486 Reproducibility

487 All the models and necessary scripts that were used in this paper can be found in the following

488 GitHub repository: https://github.com/YubinXie/multiscale-synaptic-model. Detailed guides are

<sup>489</sup> provided to reproduce all the figures.

## 490 Acknowledgements

<sup>491</sup> The authors thank Nicolas Le Novère for the discussion on the System Biology Graphical Notation

<sup>492</sup> form of biochemical model schematic. We thank Lu Li, Pınar Pir, and Varun B. Kothamachu for the

<sup>493</sup> discussion and inputs on the System Biology Markup Language version of the biochemical model.

<sup>494</sup> We also thank Sven Sahle for discussions on the COPASI software.

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