

POSTER PRESENTATION

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An improved translational switch for long term maintenance of synaptic plasticity

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Memory lasts a lifetime, yet the physiological substrate of memory, synaptic contacts, are composed of proteins that have much shorter lifetimes. A physiological analog of memory formation, long-term potentiation (LTP), has a late protein synthesis dependent phase (LLTP) that can last for many hours in slices, or even days in vivo [1,2]. Our previous studies show that maintenance of L-LTP and memory can be accounted by persistent regulation of on-site synthesis of plasticity-related proteins by a self-sustaining regulation of translation. It has been shown that a α CaMKII-CPEB1 molecular pair can act as a bistable switch with different total amounts of α CaMKII in potentiated and non-potentiated synapses [3].

The molecular interaction model in our previous study comprised a CaMKII which could be in an inactive, active and active and phosphorylated forms together with a translation regulating molecule CPEB1, which can be in an active or inactive form. The model included both degradation and new protein synthesis of αCaMKII. We have showen that this model is bistable [3]. The bistability was caused by interaction of Ca² +-Calmodulin dependent and auto-phosphorylation activation, spontaneous degradation and synthesis loops of aCaMKII. This model could successfully account for maintenance of L-LTP over a long period of time and also proposes an explanation for why application of protein synthesis and a CaMKII inhibitors at induction and maintenance phases of L-LTP result in very different outcomes [3-5]

However, the protein synthesis loop in our previous model was very simplistic. Here, we suggest a more detailed model of translation with explicit implementation of mRNA and poly-ribosome concentration in the pre-synaptic spine. We assume that activated CBEB1activates mRNA which then binds preferentially to polyribosome, as compared to a non-active mRNA, for $\alpha CaMKII$ synthesis. We show that this system can act as a bistable switch. We also look at the behavior of this system at low poly-ribosome and mRNA concentration levels using stochastic simulations with Gillespie algorithm.

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