CORRECTION

Correction: A Unifying Probabilistic View of Associative Learning

The PLOS Computational Biology Staff

There is an error in equation 15, the "discounted time derivative" h_t is defined incorrectly. It should read as follows:

"Operationally, the only change from the Kalman filter model described above is to replace the stimulus features x_t with their discounted time derivative, $h_t = x_t - \gamma x_{t+1}$. To see why this makes sense, note that the immediate reward can be expressed in terms of the difference between two values:

$$\begin{split} \mathbf{r}_t &= \mathbf{V}(\mathbf{x}_t) - \gamma \mathbf{V}(\mathbf{x}_{t+1}) \\ &= \mathbf{w}_t \mathbf{x}_t - \gamma \mathbf{w}_t \mathbf{x}_{t+1} \\ &= \mathbf{w}_t (\mathbf{x}_t - \gamma \mathbf{x}_{t+1}). \end{split} \tag{1}$$

This error does not affect the simulations, which were implemented with the correct definition.

Reference

1. Gershman SJ (2015) A Unifying Probabilistic View of Associative Learning. PLoS Comput Biol 11(11): e1004567. https://doi.org/10.1371/journal.pcbi.1004567 PMID: 26535896



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