Input Asymmetry: Left=0.1, Right=1.0

Readout Accuracy Over Training (Control Trials) - Mean ± SD across 10 seeds 1.0 Readout Accuracy (Control) 0.9 8.0 Left Hemi Readout Accuracy Right Hemi Readout Accuracy 0.7 Corruption Start (Epoch 11) 0.6 0.5 0.4 Left/Right ALM Agreement Over Training (Control Trials) - Mean ± SD across 10 seeds Empirical Agreement (L=R) 1.0 Chance Level (Binomial) Corruption Start (Epoch 11) Fraction of Trials with L=R

'0 0 0

'0 0

'0 8

'0 6

'0 8 0.5 0.4 1 2 3 4 5 6 7 8 9 10111213141516171819202122232425262728293031323334353637383940 **Epoch**