	PHASE-TTC	Garl full	Garl event-only	Garl visual-only
CCRs1-low	50/50	49/50	50/50	49/50
CCRs1-medium	43/43	42/43	43/43	42/43
CCRs1-high	30/30	29/30	30/30	29/30
CCRs2-low	46/46	45/46	46/46	45/46
CCRs2-medium	55/55	54/55	55/55	54/55
CCRs2-high	59/59	58/59	59/59	58/59
CCRm-low	42/42	41/42	42/42	41/42
CCRm-medium	40/40	39/40	40/40	39/40
Slider-750	51/51	49/51	51/51	51/51
Slider-1000	18/18	18/18	18/18	18/18

Counts show finite predictions per sequence; tinted cells mark queries with insufficient causal input.