Unifying observables through latent dynamics: shared trajectories of brain, body, and behavior

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A fundamental challenge in neuroscience is the general lack of governing laws relating macroscopic variables. With recent strides in the ability to obtain high-dimensional, multimodal recordings from awake animals, there is growing need for physiologically principled, low-dimensional descriptions of the effective dynamics. To this end, we have hypothesized the existence of an intrinsic, arousal-related dynamical process whose regulation of organism-wide physiology underpins fluctuations observed across diverse neural, physiological and behavioral measurements [1]. We support this hypothesis by introducing a computational framework that ultimately allows us to construct high-fidelity nonlinear mappings between such measurements (e.g., pupillometry and neural calcium imaging) from a latent, low-rank representation of their dynamics.

Specifically, to test our hypothesis, we introduce a data-driven approach to parsimoniously link observables that evolve according to a common but unknown dynamical mechanism. Our approach exploits Takens' embedding theorem from dynamical systems theory, which permits the (topology-preserving) reconstruction of a full-state attractor manifold from a single observable and its time history. Takens' theorem enables a strong prediction – and thus, a challenging test – of our framework: in theory, a single arousal-related observable (e.g., pupil size) should suffice to reconstruct high-dimensional observables, to the extent that a dynamical mechanism is shared. We test this prediction by performing simultaneous pupillometry and widefield calcium imaging in awake mice. The latter reports the large-scale spatial structure of cortical activity, which we hypothesize to be tightly regulated in accordance with arousal dynamics [1]. We train neural networks to approximate the hypothesized mapping from delay-embedded pupil time series (i.e., a scalar measurement) to (high-dimensional) widefield measurements, where delay embedding is achieved via dimensionality reduction of the associated Hankel matrix [2].

Following this procedure, we are able to reconstruct a surprising extent of cortex-wide spatiotemporal dynamics in awake mice from simultaneously measured pupil diameter (i.e., $74 \pm 8\%$ of variance < 0.2 Hz in held-out data, vs. $13 \pm 10\%$ without delay embedding (mean \pm std, n=7 mice)). We achieve similar success using a scalar index of behavioral arousal (whisker movement), and in predicting widefield measurements of brain metabolism or hemodynamics, suggesting the ability of our procedure to elucidate low-dimensional generative factors that underlie physiologically diverse, high-dimensional measurements. Taken together, our framework and findings elucidate an arousal-related dynamical mechanism that is both richer and substantially further-reaching than presently recognized. Our data-driven approach may be easily generalized to other settings in which a unified dynamical mechanism is hypothesized to underlie high-dimensional and potentially nonlinearly-related observables.

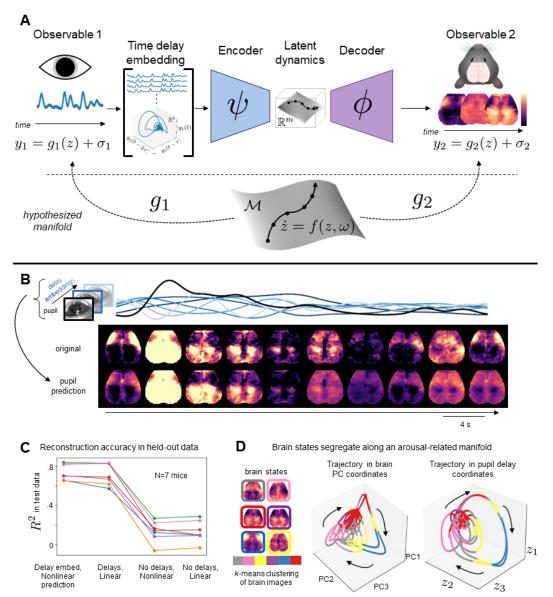


Fig. 1 (A) We consider fluctuations in diverse observables as partially reflecting a common, latent "arousal" state z and its stochastic evolution $f(z,\omega)$ occurring on a low-dimensional manifold \mathcal{M} . Formally, we represent these relations via the observation functions $g_n(z)$, each with observation-specific noise σ_n . We test the specific hypothesis that pupil size and large-scale brain activity are both tightly regulated by a common arousal-related dynamical process (cf. [1]). If true, Takens' embedding theorem implies the possibility of relating these observables via a composition of functions that map between these observables and a reconstructed latent space (i.e., ψ and ϕ). (B) Example epoch contrasting original and pupil-reconstructed widefield images (held-out data from one mouse). (C) Variance explained in held-out widefield data for each of N=7 mice. The approach described in (A) ("Delay embed, Nonlinear prediction") consistently predicts 60-85% of the observed variance < 0.2 Hz. Control analyses (see "Analysis Methods") demonstrate that delay embedding is crucial to our model's success, while estimation of ψ with a nonlinear map (via a neural network) only marginally improves performance over a multivariate linear regression model that incorporates delay embedding. (D) Example state trajectories visualized in coordinates defined by the first three widefield (brain) PCs (left) or the three leading PCs of the delay-embedded pupil time series. Trajectories are color-coded by cluster identity as determined by k-means clustering (k=6) applied to the widefield image frames. (i.e., clusters defined without reference to temporal information or pupil size).

References

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