# Dynamic sensorimotor graphs

Some Guy, Random Stranger and Anonymous Dude

Abstract—Regularities present in the somatosensory signals of a robotic agent can reflect its embodiment and the associations resulting from the active control policy. In this work, we analyze the dynamic functional connectivity of the somatosensory signals based on the instantaneous pairwise mutual information. As the robot performs exploratory motions based on motor babbling, we capture and study the timevarying changes in the signal relationships. We analyze the instantaneous and average information sharing and associate them with different information states. Results from a simulated planar system validate using the instantaneous mutual information as a tool for extracting and leveraging the relationships between the somatosensory signals and defining exploratory motions related to self-tocuh events.

#### I. INTRODUCTION

The study of the structural and functional connectivity in the brain has furthered the understanding of its organizations and information processing. An analogy with the sensorimotor signals of an embodied agent is yet to be realized. While looking at the structural connectivity of the sensorimotor signals is difficult, studying their functional connectivity to extract information about the body structure and the emergence of behaviors based on information acquisition is undoubtedly realizable. The underlying premise is that regularities among the robot's sensorimotor signals connect tightly to the robot's embodiment. These regularities, known as sensorimotor contingencies (SMC) [1], can be identified and studied using information-theoretic methods. This work explores an embodied agent's time-varying sensorimotor functional structure leveraging concepts from information and graph theory.

#### A. Related works

As established in [1], SMCs play a role in acquiring body knowledge, generalization, and goal-directedness. Yet, although the literature about sensorimotor representations is extensive [2], the connections between sensorimotor regularities and body knowledge are poorly understood. Several works [3]-[7] have used information-theoretic metrics to study these relationships in the sensorimotor system. Additionally, when exploring SMCs, touch is an essential sensory modality. For example, the work by [12] discussed using intrinsic motivation and goal-babbling for learning self-touch on a simulated humanoid robot with artificially sensitive skin. Self-touch for calibration was used in [13], letting the robot close the kinematic chain by touching its own body. In [14], a multimodal variational autoencoder in a denoising framework is used to learn the latent representation of self-touch that allows the agent to reconstruct self-reaching configurations internally.

The study of functional connectivity has shown that the statistical properties of the considered signals and their relationships may vary over time. This *dynamical* functional connectivity (DFC) has mainly been used to study functional networks in the brain to, for example, show states of lower functional connectivity during onsets of epilepsy[9] or identify potentially abnormal connectivity patterns in The work in brains affected by illness[10]. Recently it was used to identify evolving relationships in the sensorimotor signals of a simulated infant based on age [11].

#### B. Contributions

This work considers that representing the functional connections between somatosensory signals (touch and proprioception) of an embodied agent as dynamic graphs and their corresponding analysis can help understand the formation and evolution of SMCs. Given that, based on the motion policy driving the robot or the task it executes, sensorimotor connectivity will also change, capturing and studying these varying connectivity patterns is essential. We achieve this by looking at the instantaneous mutual pairwise mutual information showing how the state of information sharing in the system is related to its motion and potential interactions with itself. Using non-negative matrix factorization, we identify different information-sharing states and classify them according to their information content. In a case study, we demonstrate how, after a motor babbling phase, by only focusing on the mutual information and without knowing the robot's morphology, an excitation trajectory for both the robot arms can be devised that avoids self-contact. A comparison of this trajectory against conventional trajectory design methods is presented.

## II. THE EMBODIED AGENT

## A. The planar dual arm model

We use as reference system the robot model presented in [14], [15], consisting of a simple six degrees-of-freedom planar dual arm system with three links per arm and a fixed torso, see Fig. 2. The robot is equipped with a set of tactile sensors distributed along the robot's body. The dynamics of the model were instantiated by assigning inertial properties to the robot's composing bodies. Its actuation mechanism is based on the biology-inspired model presented in [16]–[18], where each joint is driven by antagonistic muscles (modeled as spring-damper systems) whose pulling force is linearly controlled by the signal of a corresponding motor neuron  $\sigma$ . The generated joint torque, expressed as

$$\tau = \alpha \left( \sigma_{fx} - \sigma_{ex} \right) + \beta \left( \sigma_{fx} + \sigma_{ex} + \gamma \right) q + \delta \dot{q}, \quad (1)$$

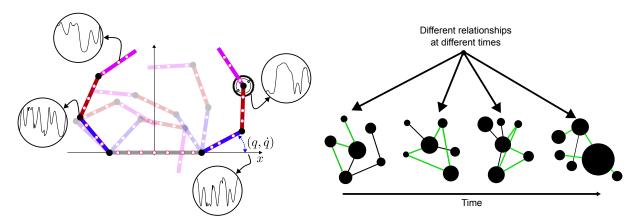


Fig. 1: General overview. While moving the sensorimotor functional connectivity changes in time

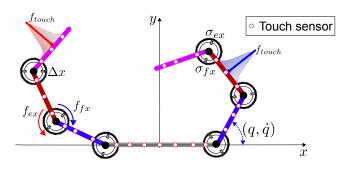


Fig. 2: The planar dual arm robot with antagonistic actuation.

results from the difference between the flexion  $\sigma_{fx}$  and extension  $\sigma_{ex}$  activation signals which create the flexion and extension pulling forces  $f_{fx}$  and  $f_{ex}$ . The parameters are:

- $\alpha$ : muscle force gain
- $\beta$ : stiffness gain
- $\gamma$ : tonic stiffness
- $\delta$ : damping coefficient

#### B. The set of somatosensory signals

The tactile sensors on the robot's body are modeled based on population coding [19] represented as distance-dependent Gaussian receptive fields (see Fig. 3), with the location of the sensors (randomly located along the robot's one-dimensional body) being the means of the receptive fields. We modified the tactile sensors to account for the strength of touch. Essentially, the previously distance-based activation of the Gaussian receptive fields is now scaled by the contact force. Similar to the tactile sensors, the robot's proprioceptive measurements are also encoded using receptive fields. Therefore, the vector of somatosensory signals s in the extended model consists of proprioception that encompasses joint position p, velocity v, and effort e, as well as touch signals r (scaled by the touch force); i.e.:

$$s = \begin{bmatrix} \boldsymbol{p}^T & \boldsymbol{v}^T & \boldsymbol{e}^T & \boldsymbol{r}^T \end{bmatrix}^T \in \mathbb{R}_{\geq}^{N_s} \tag{2}$$

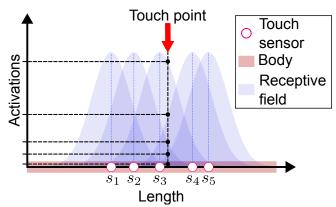


Fig. 3: The receptive fields used in population coding.

# III. THE SENSORIMOTOR DYNAMIC FUNCTIONAL CONNECTIVITY

#### A. Functional connectivity

Functional connectivity (FC) is a method for network topology inference that characterizes the dependencies of the observed signals from a system based on their probability distributions[20]. It can be subdivided into undirected and directed; the latter being related to the analysis of statistical causation from the data [21]. By studying the FC it is possible to reveal a structure that aids in the analysis of the interaction among the entities.

Based on the connection between embodiment and information structure [22], our hypothesis is that properties of the sensorimotor interactions can be made apparent by studying the FC among the somatosensory signals s(t). From the various metrics that have been proposed to evaluate such relationships, we in particular leverage those based on information theory [23], [24], as their *model-free* nature can capture linear and nonlinear relationships between signals. Particularly, we use the *mutual information* (MI), a quantity that has been applied in different contexts to quantify the relationships between variables [25]. The MI between two signals I(X, Y) can be interpreted as the amount by which

a random signal Y reduces the uncertainty about a random signal X [26]. It is a symmetric measure of the information sharing by both signals and is computed as:

$$I(X;Y) = I(Y;X) = H(X) + H(Y) - H(X,Y)$$
(3)

with the Shannon's entropy of a variable X defined by

$$H(X) = -\sum_{i=1}^{n} p(x_i) \log_2 \left( p\left(x_i\right) \right) \tag{4}$$

and the joint entropy between X and Y expressed as

$$H(X,Y) = -\sum_{i=1}^{n} \sum_{j=1}^{n} p(x_i, y_j) \log_2(p(x_i, y_j)).$$
 (5)

By extension, the mutual information matrix  $\mathcal{I} \in \mathbb{R}^{m \times m}$  can be constructed by computing the pairwise MI between the  $\left\{s_i\right\}_{i=1}^{N_s}$  somatosensory signals. Its (i,j) entries are given by

$$(\mathcal{I})_{i,j} = I(s_i, s_j). \tag{6}$$

In practice, computing (6) for a pair  $(s_i(t), s_j(t))$  involves centering their samples (to zero mean and unit standard deviation) and using either binning, kernel, or nearest neighbor methods [27] to compute their mutual information. In this work, for the computation of  $\mathcal{I}$  we use the the open-source MATLAB package *Mutual information computation*[28].

# B. Dynamic functional connectivity

When analyzing FC it might be interesting to look not only at the aggregated effect of a complete dataset of recordings but also at the instantaneous changes that occur in the relationships. Indeed, the functional relationships between sensorimotor signals can change rapidly depending on the motion policy and the interaction of the agent with the environment. To capture this time-varying, i.e. *dynamic*, functional connectivity with mutual information, it is common to use a sliding time window [29] with forward step  $\Delta t$  from which the MI is computed only for a small number of samples.

In particular, for a time window of length T. The mutual information  $I_t(s_x(t); s_y(t))$  for a distinct pair of signals  $s_x(t)$  and  $s_y(t)$  at time t is computed for the set of signal samples spanning the interval [t-T,t], see Fig. 4. We called this term the *instantaneous mutual information* (IMI). It follows that the mutual information matrix  $\mathcal{I}(t)$  at time t is constructed by computing the IMI for all the pairwise signals in the same time interval. The time series the timevarying mutual information matrix  $\mathcal{I}(t)$  shows the dynamical functional connectivity between the somatosensory signals.

#### C. Analyzing patterns

Given a set of m samples from the somatosensory signals, generated, for example, via motor babbling, a dataset  $S \in \mathbb{R}^{n \times m}$  can be used to search for repeating patterns of FC that encode certain modes of operation. It is also important to determine the strength and frequency of expression of these patterns to determine their relevance. Typical methods to detect repeating patterns in dynamic FC include the

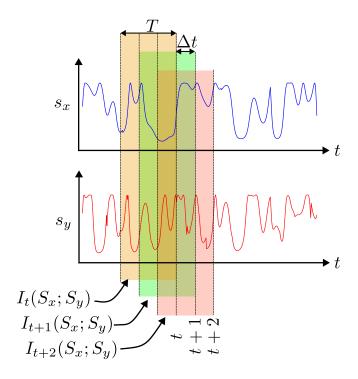


Fig. 4: Sliding window strategy to compute the instantaneous mutual information.

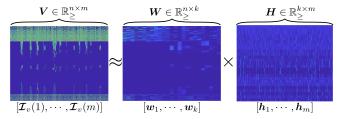


Fig. 5: Decomposition of the mutual information data matrix using non negative matrix factorization.

cosine similarity [30] and non-negative matrix factorization (NNMF) [31]. We chose the latter, as it proven useful in several studies analyzing brain dynamic functional networks and has also being used for community detection[32], [33].

NNMF is an unsupervised machine learning algorithm that can be used to split an input matrix  $D_{\mathcal{I}} \in \mathbb{R}^{n \times m}_{\geq}$  into two parts: a matrix  $W \in \mathbb{R}^{n \times k}_{\geq}$  of bases and a matrix  $H \in \mathbb{R}^{k \times m}_{\geq}$  of their corresponding contributions to the input matrix  $D_{\mathcal{I}}$ ; i.e.,

$$D_{\mathcal{I}} \approx WH.$$
 (7)

In our case, the matrix  $D_{\mathcal{I}}$  is constructed by vectorizing and concatenating each of the matrices  $\mathcal{I}_v(t) = \text{vec}(\mathcal{I}(t))$ ; that is:

$$\boldsymbol{D}_{\mathcal{T}} = [\boldsymbol{\mathcal{I}}_{n}(1), \cdots, \boldsymbol{\mathcal{I}}_{n}(m)] \tag{8}$$

Note that  $n = N_s(N_s - 1)/2$  is the total number of mutual information pairs. Since the matrix  $D_{\mathcal{I}}$  is strictly nonnegative, it is straight away suitable for its decomposition and analysis using NNMF. One crucial question is the number

of factors k used to approximate the original dataset. From the various methods to select an adequate number [34], we chose k following the elbow method as in [35] by performing NNMF for ascending values of k and selecting the value where the residual error is not reduced any further.

Each of the factors  $\{\boldsymbol{w}_i\}_i^k = 1$  can be interpreted as a base FC graph capturing a state of information sharing of multiple sensorimotor states. When the factors are aggregated using via the columns of  $\boldsymbol{H}$  the particular mutual information matrix at time t is closely approximated.

To facilitate the analysis of the relevance of each of the factor, after factorization the factors are normalized to according to the  $L_2$ -norm.

#### IV. SIMULATION RESULTS

#### A. Exploratory phase

For the computation of the instantaneous mutual information we used sensors signals sampled at 100 Hz and a sliding window of T=0.1 seconds; i.e., the previously seen 10 samples. With this short memory which is stored in a buffer, we can capture (fast) touch events.

## B. Dimensionality reduction

To show graphically how the different factors cluster naturally depending on the touched regions we used the PaCMAP dimensionality reduction method [36] given its properties to preserve aspects of the global and local structure when reducing into the latent space [37].

## V. CASE STUDY: ROBOT EXCITATION TRAJECTORIES

**TODOI**n this section we use the instantaneous mutual information to generate trajectories for the left and right arms avoiding potential collisions. This is done agnostic to the actual morphology of the robot. In contrast we use a standard method for the design of excitation trajectories and compare the results.

# VI. BEYOND ROBOTICS

# TODOValentin + Matej

# VII. CONCLUSIONS

#### REFERENCES

- [1] L. Jacquey, G. Baldassarre, V. G. Santucci, and J. K. O'Regan, "Sensorimotor contingencies as a key drive of development: From babies to robots," *Frontiers in neurorobotics*, vol. 13, p. 98, 2019.
- [2] P. D. Nguyen, Y. K. Georgie, E. Kayhan, M. Eppe, V. V. Hafner, and S. Wermter, "Sensorimotor representation learning for an "active self" in robots: A model survey," *KI-Künstliche Intelligenz*, vol. 35, no. 1, pp. 9–35, 2021.
- [3] N. M. Schmidt, M. Hoffmann, K. Nakajima, and R. Pfeifer, "Bootstrapping perception using information theory: Case studies in a quadruped robot running on different grounds," *Advances in Complex Systems*, vol. 16, no. 02n03, p. 1250078, 2013.

- [4] M. Lungarella and O. Sporns, "Mapping information flow in sensorimotor networks," *PLoS Comput Biol*, vol. 2, no. 10, e144, 2006.
- [5] D. Polani and M. Möller, "Models of information processing in the sensorimotor loop," in *Information Theory and Statistical Learning*, Springer, 2009, pp. 289–308.
- [6] T. Bossomaier, L. Barnett, M. Harré, and J. T. Lizier, "An introduction to transfer entropy," *Cham: Springer International Publishing*, vol. 65, 2016.
- [7] L. A. Olsson, C. L. Nehaniv, and D. Polani, "From unknown sensors and actuators to actions grounded in sensorimotor perceptions," *Connection Science*, vol. 18, no. 2, pp. 121–144, 2006.
- [8] X. Dong, D. Thanou, M. Rabbat, and P. Frossard, "Learning graphs from data: A signal representation perspective," *IEEE Signal Processing Magazine*, vol. 36, no. 3, pp. 44–63, 2019.
- [9] E. Christiaen, M.-G. Goossens, B. Descamps, L. E. Larsen, P. Boon, R. Raedt, and C. Vanhove, "Dynamic functional connectivity and graph theory metrics in a rat model of temporal lobe epilepsy reveal a preference for brain states with a lower functional connectivity, segregation and integration," *Neurobiology of disease*, vol. 139, p. 104 808, 2020.
- [10] T. Zhou, J. Kang, F. Cong, and X. Li, "Early childhood developmental functional connectivity of autistic brains with non-negative matrix factorization," *NeuroImage: Clinical*, vol. 26, p. 102251, 2020. [Online]. Available: https://www.sciencedirect.com/science/article/pii/S2213158220300887?via=ihub.
- [11] H. Kanazawa, Y. Yamada, K. Tanaka, M. Kawai, F. Niwa, K. Iwanaga, and Y. Kuniyoshi, "Openended movements structure sensorimotor information in early human development," *Proceedings of* the National Academy of Sciences, vol. 120, no. 1, e2209953120, 2023.
- [12] F. Gama, M. Shcherban, M. Rolf, and M. Hoffmann, "Goal-directed tactile exploration for body model learning through self-touch on a humanoid robot," *IEEE Transactions on Cognitive and Developmental Systems*, 2021.
- [13] A. Roncone, M. Hoffmann, U. Pattacini, and G. Metta, "Automatic kinematic chain calibration using artificial skin: Self-touch in the icub humanoid robot," in 2014 IEEE International Conference on Robotics and Automation (ICRA), IEEE, 2014, pp. 2305–2312.
- [14] V. Marcel, J. K. O'Regan, and M. Hoffmann, "Learning to reach to own body from spontaneous self-touch using a generative model," in 2022 IEEE International Conference on Development and Learning (ICDL), IEEE, 2022, pp. 328–335.
- [15] F. Mannella, V. G. Santucci, E. Somogyi, L. Jacquey, K. J. O'Regan, and G. Baldassarre, "Know your body through intrinsic goals," *Frontiers in neurorobotics*, p. 30, 2018.

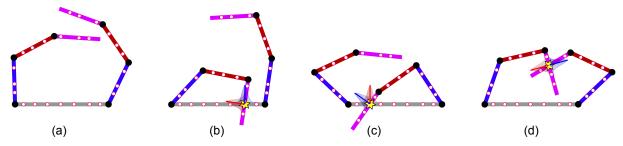


Fig. 6: Different events during exploration. (a) Pure proprioception (no touch), (b) contact with left arm, (c) contact with right arm, and (d) contact with both arms.

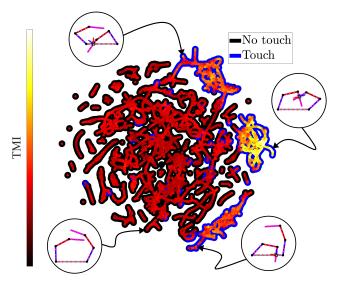


Fig. 7: A two dimensional projection using PaCMAP of the coefficients  $\mathbf{H}$ . The color of each point is scaled by its total mutual information value.

- [16] Ö. Ekeberg, "A combined neuronal and mechanical model of fish swimming," *Biological cybernetics*, vol. 69, no. 5-6, pp. 363–374, 1993.
- [17] T. Wadden and Ö. Ekeberg, "A neuro-mechanical model of legged locomotion: Single leg control," *Biological cybernetics*, vol. 79, no. 2, pp. 161–173, 1998.
- [18] Y. Shim and P. Husbands, "Chaotic exploration and learning of locomotion behaviors," *Neural computation*, vol. 24, no. 8, pp. 2185–2222, 2012.
- [19] S. Panzeri, F. Montani, G. Notaro, C. Magri, and R. S. Peterson, "Population coding," in *Analysis of Parallel Spike Trains*, S. Grün and S. Rotter, Eds. Boston, MA: Springer US, 2010, pp. 303–319, ISBN: 978-1-4419-5675-0. DOI: 10.1007/978-1-4419-5675-0\_14. [Online]. Available: https://doi.org/10.1007/978-1-4419-5675-0\_14.
- [20] K. J. Friston, "Functional and effective connectivity: A review," *Brain connectivity*, vol. 1, no. 1, pp. 13–36, 2011.
- [21] A. M. Bastos and J.-M. Schoffelen, "A tutorial review of functional connectivity analysis methods and their

- interpretational pitfalls," Frontiers in systems neuroscience, vol. 9, p. 175, 2016.
- [22] R. Pfeifer, M. Lungarella, and F. Iida, "Self-organization, embodiment, and biologically inspired robotics," *science*, vol. 318, no. 5853, pp. 1088–1093, 2007.
- [23] F. Bonsignorio, "Entropy based metrics of sensory motor coordination: A short survey," Metrics of Sensory Motor Coordination and Integration in Robots and Animals: How to Measure the Success of Bioinspired Solutions with Respect to their Natural Models, and Against More 'Artificial' Solutions?, pp. 89–110, 2020.
- [24] —, "Quantifying the evolutionary self-structuring of embodied cognitive networks," *Artificial life*, vol. 19, no. 2, pp. 267–289, 2013.
- [25] R. Steuer, J. Kurths, C. O. Daub, J. Weise, and J. Selbig, "The mutual information: Detecting and evaluating dependencies between variables," *Bioinformatics*, vol. 18, no. suppl\_2, S231–S240, 2002.
- [26] T. M. Cover, *Elements of information theory*. John Wiley & Sons, 1999.
- [27] J. Walters-Williams and Y. Li, "Estimation of mutual information: A survey," in *International Conference on Rough Sets and Knowledge Technology*, Springer, 2009, pp. 389–396.
- [28] H. Peng, Mutual information computation, https://www.mathworks.com/ matlabcentral/fileexchange/14888mutual-information-computation, Accessed: 2023-04-20.
- [29] M. G. Preti, T. A. Bolton, and D. Van De Ville, "The dynamic functional connectome: State-of-the-art and perspectives," *Neuroimage*, vol. 160, pp. 41–54, 2017.
- [30] S. S. Menon and K. Krishnamurthy, "A comparison of static and dynamic functional connectivities for identifying subjects and biological sex using intrinsic individual brain connectivity," *Scientific reports*, vol. 9, no. 1, p. 5729, 2019.
- [31] X. Fu, K. Huang, N. D. Sidiropoulos, and W.-K. Ma, "Nonnegative matrix factorization for signal and data analytics: Identifiability, algorithms, and applications.," *IEEE Signal Process. Mag.*, vol. 36, no. 2, pp. 59–80, 2019.

- [32] F. Wang, T. Li, X. Wang, S. Zhu, and C. Ding, "Community discovery using nonnegative matrix factorization," *Data Mining and Knowledge Discovery*, vol. 22, pp. 493–521, 2011.
- [33] X. Luo, Z. Liu, L. Jin, Y. Zhou, and M. Zhou, "Symmetric nonnegative matrix factorization-based community detection models and their convergence analysis," *IEEE Transactions on Neural Networks and Learning Systems*, vol. 33, no. 3, pp. 1203–1215, 2021.
- [34] L. Muzzarelli, S. Weis, S. B. Eickhoff, and K. R. Patil, "Rank selection in non-negative matrix factorization: Systematic comparison and a new mad metric," in 2019 International Joint Conference on Neural Networks (IJCNN), IEEE, 2019, pp. 1–8.
- [35] H. Phalen, B. A. Coffman, A. Ghuman, E. Sejdić, and D. F. Salisbury, "Non-negative matrix factorization reveals resting-state cortical alpha network abnormalities in the first-episode schizophrenia spectrum," *Biological Psychiatry: Cognitive Neuroscience and Neuroimaging*, vol. 5, no. 10, pp. 961–970, 2020.
- [36] Y. Wang, H. Huang, C. Rudin, and Y. Shaposhnik, "Understanding how dimension reduction tools work: An empirical approach to deciphering t-sne, umap, trimap, and pacmap for data visualization," *The Journal of Machine Learning Research*, vol. 22, no. 1, pp. 9129–9201, 2021.
- [37] H. Huang, Y. Wang, C. Rudin, and E. P. Browne, "Towards a comprehensive evaluation of dimension reduction methods for transcriptomic data visualization," *Communications biology*, vol. 5, no. 1, p. 719, 2022.