Hierarchical Frequency Tagging Probe (HFTP): A Unified Approach to Investigate Syntactic Structure Representations in Large Language Models and the Human Brain

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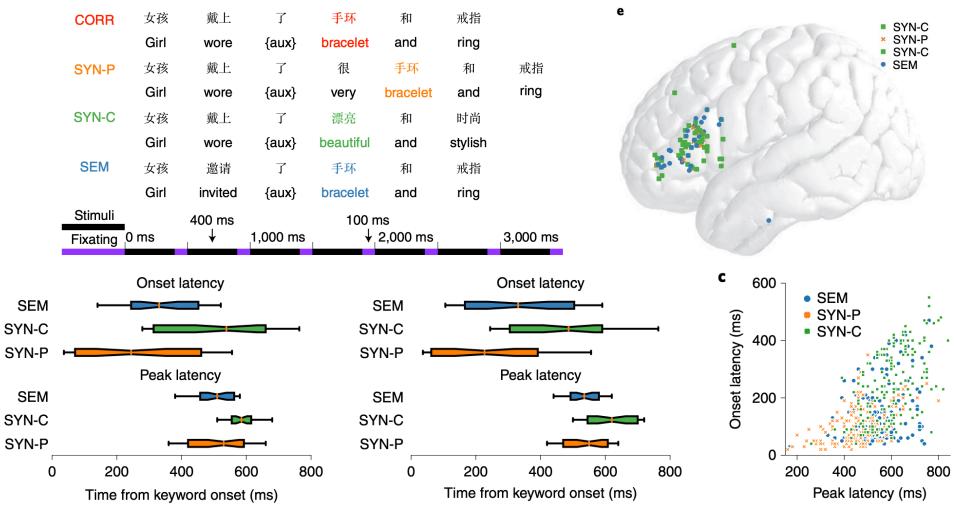
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Background - Syntactic study in the human brain



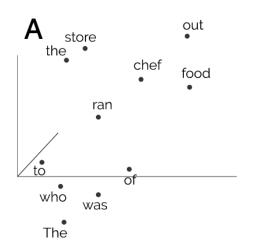


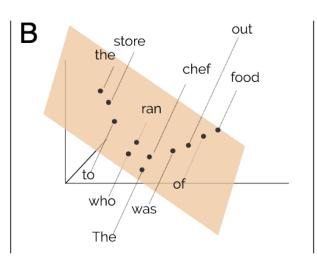
• Using syntactic-violation and semantic paradigms, single-electrode analyses in the **IFG** indicate that, **Chinese syntax is** processed before semantics; at larger scales, however, the two are intertwined.

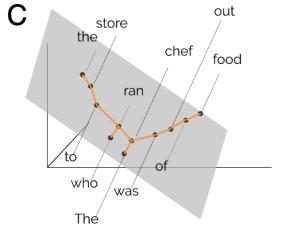


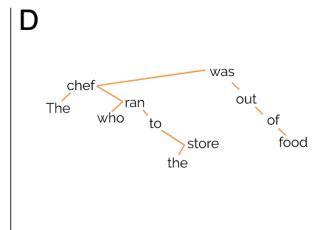
Background - Structural probe in language models











Each of the words of the sentence *The* chef who ran to the store was out of food is internally represented in context as a vector.

A structural probe finds a linear transform of that space under which squared L_2 distance between vectors best reconstructs tree path distance between words.

Once in this latent space, the structure of the tree is globally represented by the geometry of the vector space, meaning words that are close in the space are close in the tree.

In fact, the tree can be approximately recovered by taking a minimum spanning tree in the latent syntax space.

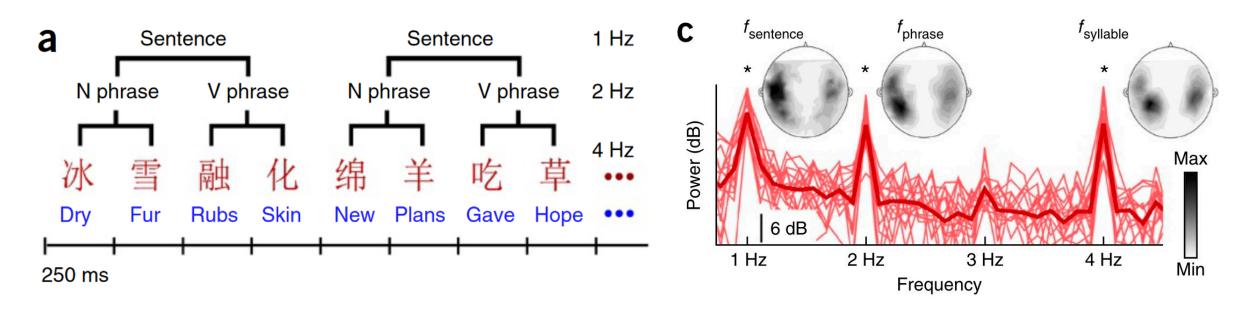
A structural probe linearly projects language-model vectors so distances approximate parse paths,
 enabling a minimum spanning tree to recover syntax.

Manning, C. D. et al. Emergent linguistic structure in artificial neural networks trained by self-supervision. PNAS (2020).



Background - Hierarchical Frequency Tagging (HFT)



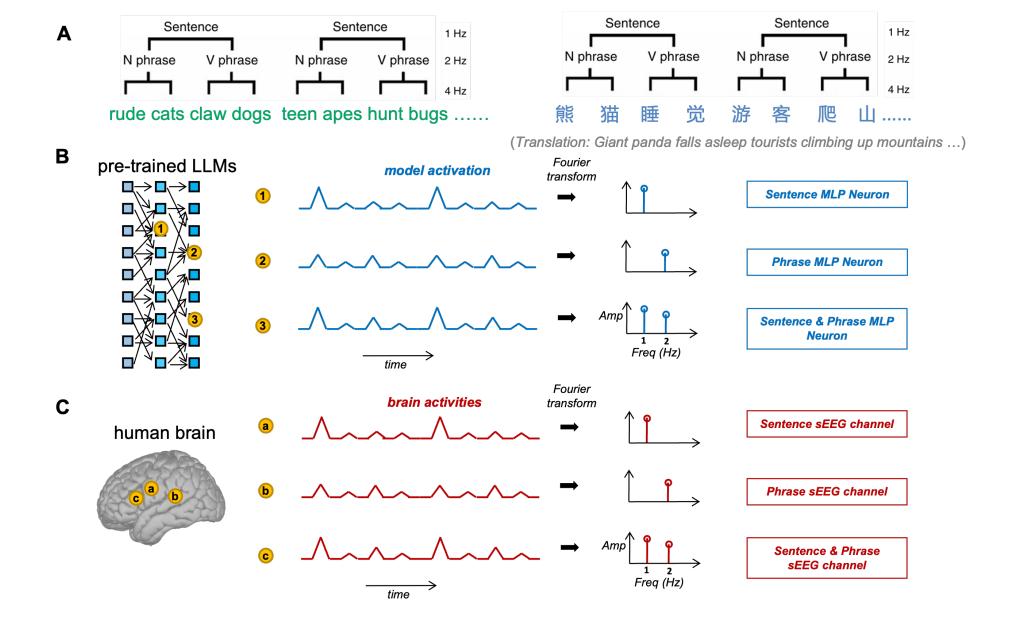


• For every four-word collocation, participants can intrinsically encode four syllables (4Hz), two phrases (2Hz) and one sentence (1Hz), and this hierarchical pattern tracking generalizes across languages.



Workflow

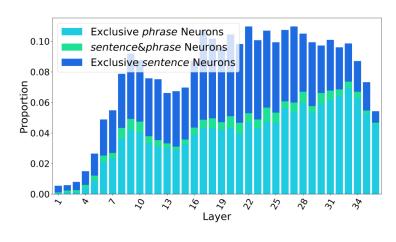






Results - Distribution of syntactic neurons

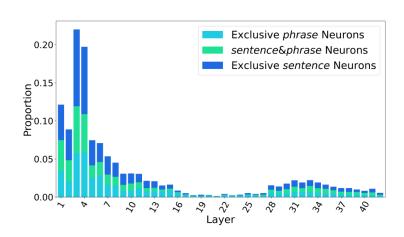




Exclusive phrase Neurons
sentence&phrase Neurons
Exclusive sentence Neurons

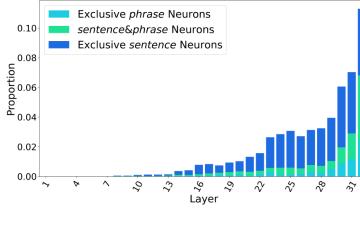
0.20
0.10
0.05
0.00

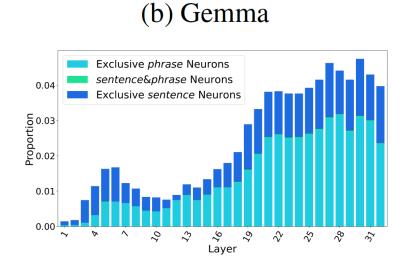
Layer

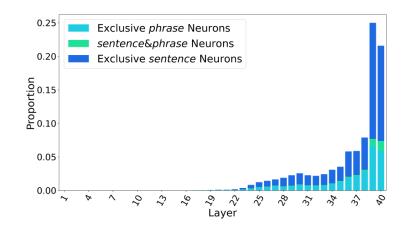


(a) GPT-2

(c) Gemma 2







(d) Llama 2

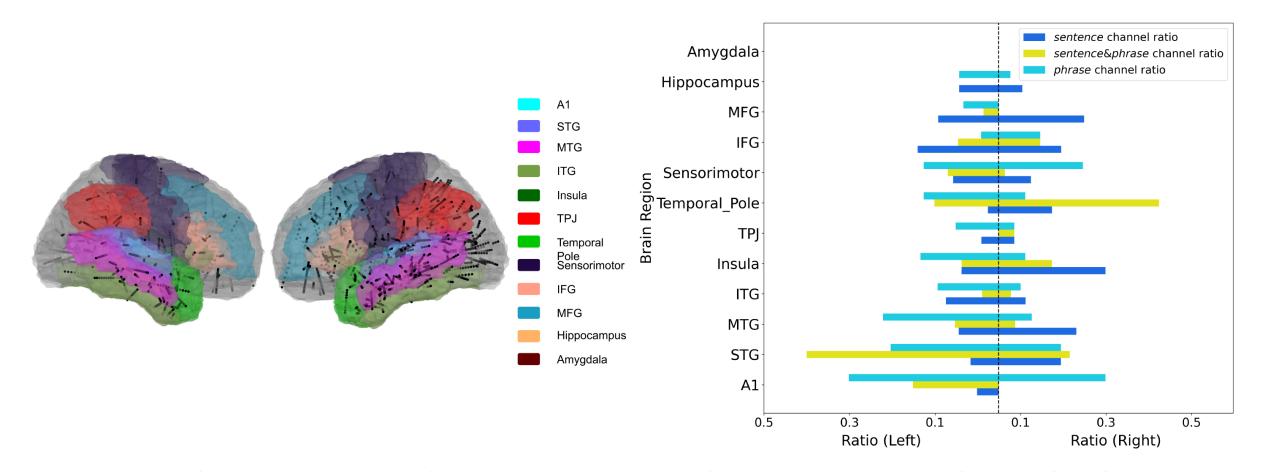
(e) Llama 3.1

(f) GLM-4



Results - Distribution of syntactic channels and Brain ROIs





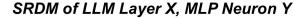
(a) sEEG channel locations and Brain ROIs

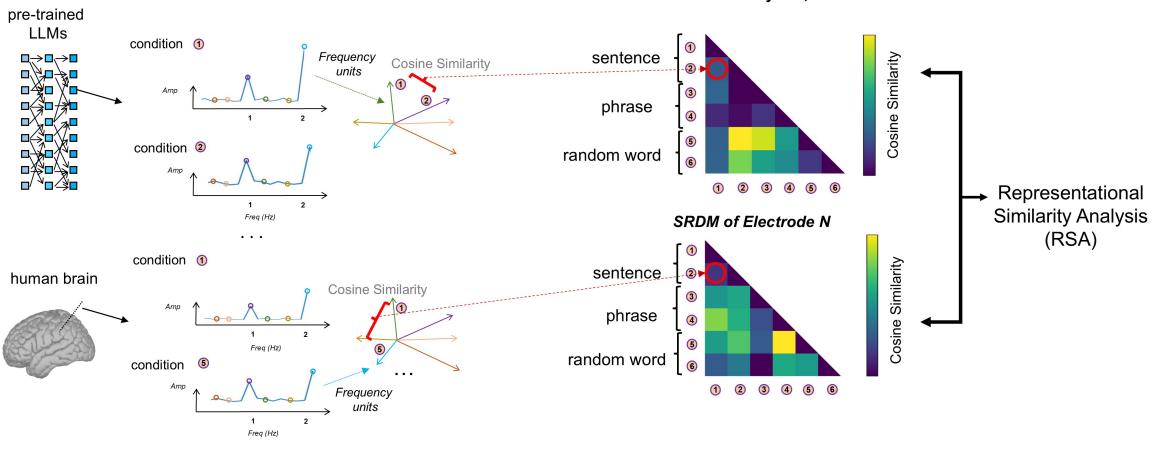
(b) Significant sEEG channel distribution



Alignment pipeline







We compute Structure Representational Dissimilarity Matrixes (SRDMs) from cosine similarities across conditions
for sentence, phrase units in LLMs and the human brain, then apply Representational Similarity Analysis (RSA)
between model and brain SRDMs to quantify model—brain correspondence.



Results – Representational Alignment



	GPT-2		Gemma		Gemma 2		Llama 2		Llama 3.1		GLM-4	
	L	R	L	R	L	R	L	R	L	R	L	R
S(m,b)	0.654	0.442	0.582	0.411	0.644	0.450	0.645	0.439	0.514	0.405	0.630	0.445
A1	0.683	0.423	0.642	0.358	0.702	0.333	0.649	0.547	0.514	0.403	0.664	0.374
STG	0.667	0.422	0.593	0.386	0.654	0.410	0.672	0.453	0.507	0.392	0.647	0.412
MTG	0.674	0.392	0.584	0.383	0.659	0.411	0.67 4	0.409	0.521	0.408	0.645	0.397
ITG	0.637	0.444	0.578	0.406	0.631	0.448	0.629	0.426	0.509	0.401	0.615	0.439
Insula	0.624	0.460	0.551	0.425	0.600	0.476	0.630	0.446	0.518	0.422	0.604	0.475
TPJ	0.610	0.452	0.566	0.373	0.641	0.410	0.619	0.400	0.518	0.408	0.606	0.438
Temporal Pole	0.648	0.473	0.556	0.470	0.643	0.558	0.610	0.469	0.494	0.448	0.616	0.483
Sensorimotor	0.637	0.462	0.567	0.426	0.622	0.448	0.624	0.446	0.505	0.396	0.617	0.463
IFG	0.694	0.463	0.603	0.466	0.670	0.496	0.665	0.491	0.513	0.410	0.646	0.490
MFG	0.615	0.436	0.557	0.401	0.585	0.489	0.597	0.367	0.510	0.397	0.588	0.473
Hippocampus	0.698	0.405	0.553	0.408	0.626	0.428	0.657	0.413	0.534	0.390	0.613	0.434
Amygdala	/	0.489	0.566	0.454	/	0.472	/	0.558	0.496	0.377	/	0.508

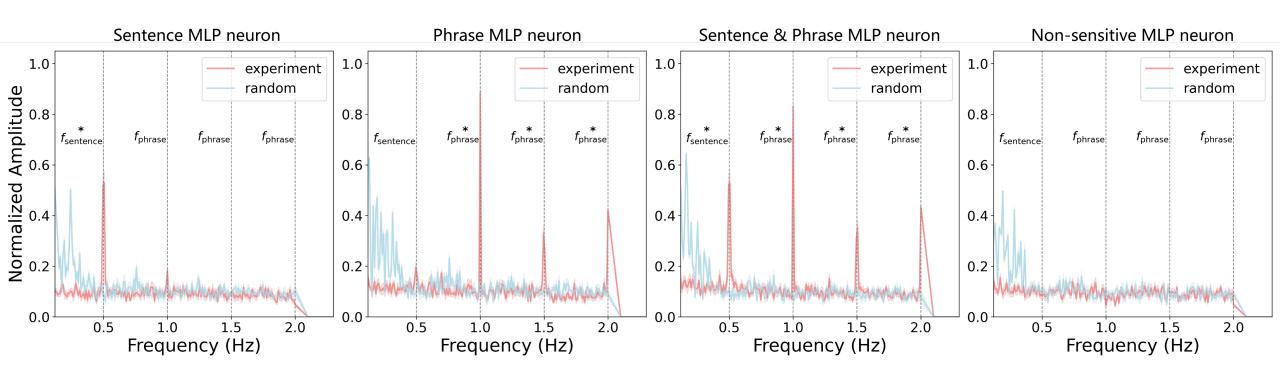
^{*} **sentence** corpus

- GPT-2 showed the strongest alignment in left hemisphere.
- Advancing models diverge in model-brain alignment (e.g., Gemma 2 ↑, Llama 3.1 ↓), uncoupling performance gains from syntactic brain-likeness.



Results – HFTP on naturalistic corpus (8-word)



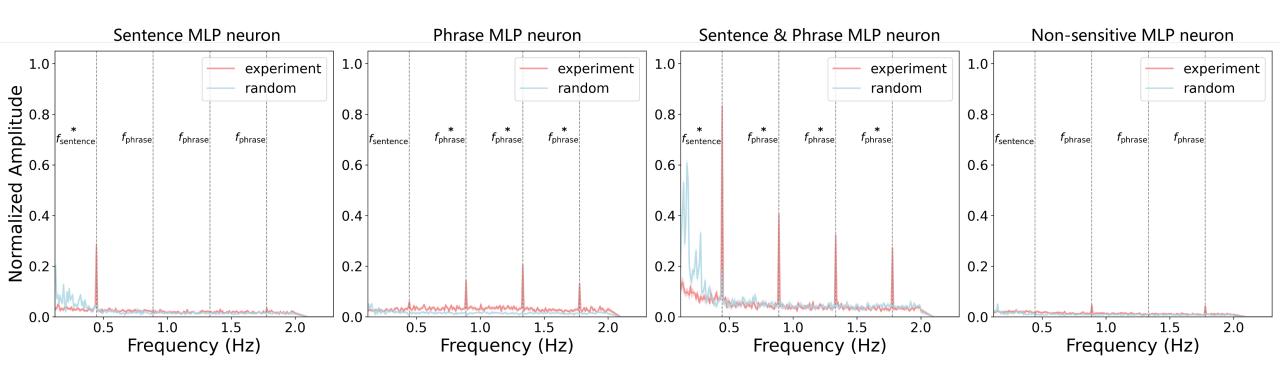


• In the 8-word English naturalistic corpus, LLMs exhibit robust peaks at the sentence rate (0.5 Hz) and at phrase rates (1, 1.5, 2 Hz), demonstrating the generalizability of our method.



Results – HFTP on naturalistic corpus (9-word)





• In the 9-word English naturalistic corpus, LLMs exhibit robust peaks at the sentence rate (~0.44 Hz) and at phrase rates (~0.89, 1.33, 1.78 Hz), demonstrating the generalizability of our method.

Conclusion



- We introduced the **Hierarchical Frequency-Tagging Probe (HFTP)**, a unified framework that **probes** internal representational structure and systematically assesses the alignment of syntactic representations between LLMs and the human brain.
- For LLMs, syntactic units cluster at different layers across models: some concentrate in **early layers** while some cluster in **later layers**.
- For the human brain, sentence- and phrase-selective sEEG channels cluster in left hemisphere—A1,
 STG, MTG, IFG—with fewer right-hemisphere.
- Representational alignment is strongest in left-lateralized language regions and varies by model family;
 upgrades do not monotonically improve brain alignment.