## 01-Groq.com API for Dev set

In this notebook, we will make API calls to Grog.com to summarize the article (up to 20k characters) in the mini (10% random sampling) dev set.

We would use Mixtral 8x7B as it has very good performance scores, and it's open source.

```
In [1]: # !pip install groq
In [2]: # Load api key
        import json
        import urllib.parse
        with open('./data/credentials.json') as f:
            login = json.load(f)
        api_key = login["GROQ_API_KEY"]
        print(len(api_key))
       56
In [3]: from groq import Groq
        client = Groq(
            api_key=api_key,
In [4]: completion = client.chat.completions.create(
            model="mixtral-8x7b-32768",
            messages=[
                     "role": "user",
                     "content": "Answer in 1 sentence: What's the capital city of Alberta?"
                },
            ],
            temperature=1,
            max_tokens=1024,
            top_p=1,
            stream=True,
            stop=None,
        for chunk in completion:
            print(chunk.choices[0].delta.content or "", end="")
```

The capital city of Alberta is Edmonton.

```
In [5]: # test some summarization
        s = 'The evolutionary origins of the hypoxia-sensitive cells that trigger amniote r
        s
```

Out[5]: 'The evolutionary origins of the hypoxia-sensitive cells that trigger amniote resp iratory reflexes – carotid body glomus cells , and 'pulmonary neuroendocrine cell s' ( PNECs ) – are obscure . Homology has been proposed between glomus cells , whi ch are neural crest-derived , and the hypoxia-sensitive 'neuroepithelial cells' ( NECs ) of fish gills , whose embryonic origin is unknown . NECs have also been lik ened to PNECs , which differentiate in situ within lung airway epithelia . Using g enetic lineage-tracing and neural crest-deficient mutants in zebrafish , and physi cal fate-mapping in frog and lamprey , we find that NECs are not neural crest-deri ved , but endoderm-derived , like PNECs , whose endodermal origin we confirm . We discover neural crest-derived catecholaminergic cells associated with zebrafish ph aryngeal arch blood vessels , and propose a new model for amniote hypoxia-sensitive cell evolution: endoderm-derived NECs were retained as PNECs , while the carotid body evolved via the aggregation of neural crest-derived catecholaminergic ( chrom affin ) cells already associated with blood vessels in anamniote pharyngeal arches . '

```
discover neural crest-derived catecholaminergic cells associated with zebrafish ph
        e cell evolution: endoderm-derived NECs were retained as PNECs , while the carotid
        body evolved via the aggregation of neural crest-derived catecholaminergic ( chrom
In [6]: prompt = f"[INST] Simplify and summarize in 200 to 300 words: {s} [/INST]"
In [7]: # test pause
        import time
        print("Hello")
        time.sleep(2)
        print("World")
       Hello
       World
In [8]: SLEEP_TIME = 10 # pause between requests
In [9]: # put into a function
        def send_sumarize_request(content, model=client, min_words=250, max_words=500, quie
                summarize the content
                input: context (text), model (groq_api client), max_words (int)
                output: summarized text
            if not quiet:
                print("Sending request for text =", content[:100])
            prompt = f'Simplify and summarize in minimum {min_words} to maximum {max_words}
            try:
                completion = client.chat.completions.create(
                    model="mixtral-8x7b-32768",
                    messages=[
                         {
                             "role": "user",
                             "content": prompt
                        },
                    ],
                    temperature=0.8,
                    max_tokens=2048,
                    top_p=1,
                    stream=True,
                    stop=None,
```

```
for chunk in completion:
    result += chunk.choices[0].delta.content or ""

except Exception as err:
    print("Skipping, error : ", err)
    result = ""

# pause to avoid hitting bandwidth limit (~ 14K token / minute)
print(f"Pausing for {SLEEP_TIME} secs...")
time.sleep(SLEEP_TIME)
print("OK")

return result
```

```
In [10]: send_sumarize_request(s)
```

Pausing for 10 secs...
OK

Out[10]: 'The evolutionary origins of hypoxia-sensitive cells responsible for amniote respi ratory reflexes, specifically carotid body glomus cells and pulmonary neuroendocri ne cells (PNECs), are unclear. Although glomus cells are neural crest-derived, it has been suggested that they share a common origin with hypoxia-sensitive neuroepi thelial cells (NECs) found in fish gills, whose embryonic origin is unknown. NECs have also been compared to PNECs, which differentiate in situ within lung airway e pithelia. However, through the use of genetic lineage-tracing, neural crest-defici ent mutants in zebrafish, and physical fate-mapping in frog and lamprey, it has be en determined that NECs are not neural crest-derived, but instead, endoderm-derive d, like PNECs, whose endodermal origin has now been confirmed. The study also disc overed neural crest-derived catecholaminergic cells associated with zebrafish phar yngeal arch blood vessels. Therefore, the study proposes a new model for the evolu tion of amniote hypoxia-sensitive cells: endoderm-derived NECs were retained as PN ECs, while the carotid body evolved via the aggregation of neural crest-derived ca techolaminergic (chromaffin) cells already associated with blood vessels in anamni ote pharyngeal arches.'

```
In [11]: import pandas as pd
```

Out[12]:		lay_summary	article	headings	keywords	id			
	0	It can take several months , or even years , f	Mature neural networks synchronize and integra	[Abstract, Introduction, Results, Discussion,	[neuroscience]	elife- 69011- v2			
	1	Many of our decisions are made on the basis of	Many decisions are thought to arise via the ac	[Abstract, Introduction, Results, Discussion,	[neuroscience]	elife- 17688- v1			
	2	Oculo-Cerebro- Renal syndrome of Lowe ( Lowe sy	Mutations in the inositol 5- phosphatase OCRL c	[Abstract, Introduction, Results, Discussion,	[cell biology]	elife- 02975- v2			
	3	When an embryo develops , its cells must work	Gradients of signaling proteins are essential	[Abstract, Introduction, Results, Discussion,	[developmental biology]	elife- 38137- v3			
	4	Our genomes contain a record of historical eve	Similarity between two individuals in the comb	[Abstract, Introduction, Results, Discussion,	[evolutionary biology, genetics and genomics]	elife- 15266- v1			
In [13]:	rar k =	port random ndom.seed(42) = random.randint(0 em = df.iloc[k] em	, len(df) - 1)						
Out[13]:	he ke id	lay_summary Spoken language is colored by fluctuations in  article Individuals with congenital amusia have a life  headings [Abstract, Introduction, Results, Discussion,  keywords [neuroscience]  id elife-53539-v2  Name: 20, dtype: object							
In [14]:	ite	em.article							

Out[14]: "Individuals with congenital amusia have a lifelong history of unreliable pitch pr ocessing . Accordingly , they downweight pitch cues during speech perception and i nstead rely on other dimensions such as duration . We investigated the neural basi s for this strategy. During fMRI, individuals with amusia (N = 15) and control s ( N = 15 ) read sentences where a comma indicated a grammatical phrase boundary . They then heard two sentences spoken that differed only in pitch and/or duration cues and selected the best match for the written sentence . Prominent reductions i n functional connectivity were detected in the amusia group between left prefronta l language-related regions and right hemisphere pitch-related regions , which refl ected the between-group differences in cue weights in the same groups of listeners . Connectivity differences between these regions were not present during a control task . Our results indicate that the reliability of perceptual dimensions is linke d with functional connectivity between frontal and perceptual regions and suggest a compensatory mechanism . \n Congenital amusia is a rare condition characterized by impaired perception of and memory for pitch ( Peretz et al . , 2002 ) . Althoug h congenital amusia presents as an auditory condition , auditory cortical response s are normal ( Moreau et al . , 2013; Norman-Haignere et al . , 2016 ) , as is sub cortical encoding of pitch ( Liu et al . , 2015b ) . The dominant view of amusia's neural basis is that connectivity between right inferior frontal cortex and right auditory cortex is impaired , resulting in impaired conscious access to pitch info rmation for guiding behavior ( Hyde et al . , 2011; Albouy et al . , 2013; Leveque et al . , 2016; Zendel et al . , 2015; see Peretz , 2016 for review ) . While cong enital amusia is believed to be innate , there is evidence that recovery is possib le through training ( Whiteford and Oxenham , 2018 ) . Although pitch is usually a ssociated with music , it is also important for cueing categories in spoken langua ge ( de Pijper and Sanderman , 1994; Streeter , 1978 ) and conveying emotion in sp eech (Frick , 1985 ) . In highly controlled laboratory tasks in which speech perc eption judgments must be made based on pitch alone , only minor deficits have been observed in amusia ( Liu et al . , 2015a; Patel et al . , 2008 ) . In naturalistic speech perception contexts , people with amusia rarely report any difficulties ( L iu et al . , 2010 ) . This may be because , in natural speech , pitch variation te nds to co-occur with variation in other acoustic dimensions , such as duration and amplitude . Our lab has shown that in such cases where multiple redundant cues are available , English-speaking individuals with amusia tend to rely less on pitch th an non-amusic controls , suggesting they may calibrate their perception by down-we ighting the cues that are less reliable for them ( Jasmin et al . , 2020a ) . As f or emotional prosody in speech , individuals with amusia can recognize emotions in spoken sentences , but not in short samples such as isolated vowels ( Pralus et al . , 2019 ) , or when speech has been filtered to remove high-frequency non-pitch c ues ( Lolli et al . , 2015 ) . It is unknown how decreased reliance on a particula r acoustic cue during speech perception ( such as pitch cues in amusia ) is reflec ted in the brain . Previous neural studies of cue integration have focused on inte gration of multiple modalities , for\xa0example the 'weighted connections' model o f multisensory integration . In this model , the relative reliability of the modal ities involved with perception of a stimulus is related to differential connectivi ty strength ( Beauchamp et al . , 2010; Rohe and Noppeney , 2018 ) . For example , when participants simultaneously view and feel touches to the hand , and reliabili ty of visual and tactile perception is manipulated experimentally via introduction of noise , connection strength ( effective connectivity measured with functional M RI and structural equation modeling ) between unimodal and multimodal sensory area s adjusts accordingly . More concretely , when visual information is degraded , th e connection strength between lateral occipital cortex ( a visual area ) and intra parietal sulcus ( a multimodal area ) decreases , and when tactile perception is m ade noisier, connection strength between secondary somatosensory cortex and intra parietal sulcus becomes weaker ( Beauchamp et al . , 2010 ) . Similarly , effectiv e connectivity between the ( multimodal ) superior temporal sulcus ( STS ) and vis

ual and auditory areas has shown similar modulations during processing of audiovis ual speech: connection strength between auditory cortex and the STS is weaker when noise has been introduced to the auditory speech , and conversely connection stren gth between visual cortex and STS is weaker if visual noise is introduced ( Nath a nd Beauchamp , 2011 ) . Just as connectivity differences have been shown to reflec t the precision of different sensory modalities during multisensory integration , an analogous phenomenon may be at work within a single modality during multidimens ional integration . As mentioned , the acoustic speech signal carries multiple cooccurring acoustic dimensions ( e . g . roughly described as voice pitch , duratio n , and amplitude ) , which often provide redundant cues to disambiguate a linguis tic category ( Patel , 2014; Winter , 2014; Jasmin et al . , 2020a ) . Individuals with typical pitch perception have learned through a lifetime of experience with s peech acoustics that vocal pitch is a useful and reliable cue . By contrast , indi viduals with amusia , who have unreliable perception of and memory for pitch ( ana logous to the 'noise' introduced in the multisensory integration studies cited abo ve ) , would have learned that , for them , pitch is not a reliable cue for proces sing spoken language . Thus , by analogy to the multisensory weighting results des cribed above , we hypothesize that amusics may exhibit decreased connectivity betw een language regions and pitch-related areas during speech processing . The neural foundations of perceptual weighting in speech have thus far not been investigated in atypical individuals . Indeed , only one previous functional neuroimaging study has examined the neural processing of spoken material in people with amusia . In t his study , no group differences were detected in task-related activation or funct ional connectivity during processing of speech ( whereas group differences were ob served during processing of tones; Albouy et al . , 2019 ) . However , the connect  $% \left( 1\right) =\left( 1\right) \left( 1\right) +\left( 1\right) \left( 1\right) \left( 1\right) +\left( 1\right) \left( 1\right) \left($ ivity analyses in this study focused on the silent retention interval in a task in which participants needed to maintain phonemic and not pitch-related information i n memory; the analyses also used broader bilateral ROIs within networks associated with language processing . It remains an open question how functional connectivity in amusic and non-amusic participants may differ during speech encoding in pitch-r elated language tasks within regions of interest selected with a whole-brain datadriven approach. To determine whether the relative reliability of auditory dimens ions in speech perception is reflected in functional connectivity , we used functi onal magnetic resonance imaging to scan 15 individuals with amusia and 15 controls . Participants matched spoken sentences with visually presented ones on the basis of the position of intonational phrase boundaries . These intonation changes were conveyed differently , in three conditions: Pitch-Informative ( where only pitch c ues could be used to make the judgment ) , Duration-Informative ( where only durat ion cues could be used ) or Both-Informative ( both pitch and duration cues could be used; Jasmin et al . , 2020a; Jasmin et al . , 2020b ) . Functional connectivit y was then examined using a data-driven approach that allowed us to identify the l argest group differences , without the need for regions of interest to be selected a priori . The benefit of this approach is that any set of regions could emerge , not only ones reported in previous literature . Crucially , task performance was m atched between the groups (based on prior behavioural testing; Jasmin et al., 2 020a ) , ensuring that any neural differences did not simply represent an inabilit y to perform the task . Finally , functional connectivity between these areas was analyzed with respect to prosodic cue weights obtained outside the scanner, and a lso compared to functional connectivity calculated from different scanning runs wi th a passive listening task . \n On each trial , participants read one visually pr esented text sentence , then heard two auditory versions of the sentence , only on e of which contained an acoustically conveyed phrase boundary in the same place as in the text sentence ( see Figure 1 for schematic and example sentences ) . Trials were scored as correct if a participant pressed the button associated with the aud itory sentence that correctly matched the text sentence . Proportions of correct j udgments (Figure 2 ) were subjected to a repeated-measures analysis of variance .

Overall , proportion correct across amusia and control groups was matched ( main e ffect of Group , F ( 1 , 84 ) = 0 . 16 , p=0 . 69 , interaction of Group by Condit ion , F ( 2 , 84 ) = 0 . 374 , p=0 . 96 ) . This lack of interaction was predicted based on previous results obtained from a similar paradigm using out-of-scanner da ta but from the same participants ( Jasmin et al . , 2020a ) . There was a main ef fect of condition ( F ( 2 , 84 ) = 3 . 32 , p=0 . 04 ) . Follow-up post-hoc testin g indicated that performance in the Both-Informative condition ( with pitch and du ration cues simultaneously present ) was more accurate than either Pitch-Informati ve ( t ( 84 ) = 2 . 31 , p=0 . 023 ) or Duration-Informative ( t ( 84 ) = 2 . 15 , p=0 . 03 ) , a result that was also predicted and which replicates the behavioral findings in Jasmin et al . , 2020a . One outlier control participant's performance was less than 0 . 3 . Re-analysis of the data without this participant did not cha nge the results pattern . Results from these analyses are available online ( see D ata Availability Statement for details ) . A data-driven approach was taken to ide ntify brain regions with the largest group- and condition-related differences in f unctional connectivity ( see Materials\xa0and\xa0methods ) . Comparing whole-brain connectedness values by group ( Amusia vs . Controls ) revealed four significant 1 ocations ( where z of peak vertices $\xa0$  $\xa0$ 4 . 61 , FDR-corrected p<0 . 05 ) that showed greater whole-brain connectedness for the control than for the amusia group ( see Figure 3 , yellow crosses ) . All group differences were located in the infe rior frontal cortex: two left hemisphere vertices (inferior frontal gyrus p . tri angularis and dorsolateral prefrontal cortex ) and two right hemisphere vertices ( inferior frontal gyrus p . triangularis and p . orbitalis ) . There were no areas where whole-brain connectedness differed by Condition , or showed an interaction o f Group and Condition . Follow-up testing was conducted on the four significant re gions ( Control\xa0>\xa0Amusia , collapsed across the three conditions ) identifie d above to characterize the specific cortical regions driving these group connecti vity differences ( Berman et al . , 2016; Gotts et al . , 2012; Jasmin et al . , 2 019; Song et al . , 2015 ) . Relative to control participants , amusic participant s' left inferior frontal gyrus seed region showed particularly notable decreases i n connectivity with the right posterior superior temporal and inferior parietal co rtex , as well as with the right posterior superior temporal sulcus ( Figure 3A ) . Analysis of subcortical connectivity indicated that there was also weaker connec tivity with the right nucleus accumbens ( Table 1 ) . The left dorsolateral prefro ntal cortex in amusic participants showed decreased functional connectivity with t he mid portions of the right superior temporal gyrus , posterior part of the right middle temporal gyrus extending into the inferior bank of the superior temporal su lcus , and the right anterior insula (Figure 3A ) . Several subcortical structure s - bilateral caudate nucleus and putamen , bilateral pallidum , bilateral cerebel lum , and bilateral thalamus - also showed significantly reduced ( FDR-corrected ) connectivity with the seed in amusics ( Table 1 ) . The right pars triangularis se ed showed Control\xa0>\xa0Amusic connectivity with right dorsolateral prefrontal c ortex and left posterior superior temporal gyrus (Figure 3B). It also showed de creased connectivity with left nucleus accumbens . Right pars orbitalis showed dec reased connectivity with right dorsolateral prefrontal cortex ( Figure 3B ) . Ther e was also decreased connectivity with the left thalamus ( Table 1 ) . Of the 30 p articipants in this study , 21 took part in an experiment that measured the degree to which they relied on pitch versus duration to categorize prosody , that\xa0is , their 'normalized prosodic cue weights', which ranged from 0 to 1 , with values  $\ensuremath{\mathbf{g}}$ reater than 0 . 5 indicating greater reliance on pitch than duration , and values less than 0 . 5 indicating greater reliance on duration than pitch ( Experiment 1 , Jasmin et al . , 2020a ) . These cue weights were assessed with respect to the f unctional connectivity results reported above . Across this subset of participants , normalized cue weights were correlated with L-DLPFC\xa0<=>\xa0R insula connectiv ity ( Spearman R\xa0=\xa00  $\cdot$  78 , p=0  $\cdot$  000037 ) , and L-DLPFC\xa0<=>\xa0R auditor y\xa0cortex connectivity ( Spearman R\xa0=\xa00 . 75 , p=0 . 000154; Figure 4 ) .

This indicated that participants who relied least on pitch information to process speech had the weakest functional connectivity between these areas , while those w ho relied most on pitch had the strongest . Although analyzing the control and amu sic groups independently results in extremely small sample sizes , this pattern al so held ( albeit with 'marginal significance' ) within the 11 control participants alone , for both auditory cortex connectivity ( R\xa0=\xa00 . 58 , p=0 . 06 ) and insular connectivity (  $R \times 0 = x_0 \cdot 55$  ,  $p=0 \cdot 08$  ) . Both these correlations were e in the predicted direction , suggesting that even non-amusics may perform dimens ional reweighting of acoustic dimensions and functional connectivity . Correlation s within the ( much more variable ) amusic group alone were weaker and non-signifi cant ( although again , the group size is very small ) . To ensure that\xa0the pat tern of connectivity we observed between groups ( decreased right auditory cortex and right insula with L-DLPFC connectivity ) was not due to intrinsic , task-irrel evant differences in neural architecture , the data from the language task was com pared to that collected during passive listening to tone sequences . Whereas durin g speech perception amusic subjects showed reduced functional connectivity between left frontal and right insula/auditory ROIs relative to controls ( p=0 . 0001 for both ROIs; in line with the whole-brain imaging analyses ) , this pattern did not hold during passive listening to tones ( Amusia vs Control connectivity , p=0 . 29 , Group ( Amusic , Control ) by Task ( Speech Perception , Passive Tone Listening ) interaction p=0 . 045 for the insula ROI; Amusia vs Control p=0 . 30 , Group by Task interaction p=0 . 035 for the auditory cortex ROI - see Figure 5 ) . These in teractions suggest that our neural connectivity results are specifically linked to speech perception , rather than reflecting an overall connectivity difference betw een groups regardless of task state . Although we were concerned with functional c onnectivity rather than activation , we also tested for differences in activation levels between groups and conditions . False Discovery Rate correction was used to correct for multiple comparisons across both hemispheres for each test ( Group , C ondition and Group X Condition ) . No significant differences were detected for th e main effects of group and condition , nor the interaction of those factors . \n We found that individuals with amusia , who have been previously shown to rely les s on pitch than controls to process spoken language ( Jasmin et al . , 2020a ) , e xhibited decreased functional connectivity between left frontal areas and right he misphere pitch-related regions . In our task , participants matched spoken sentenc es with visually presented sentences based on pitch , duration , or both these aco ustic dimensions together . Using a data-driven approach , we identified four regi ons in left and right inferior frontal cortex for which the amusic group exhibited decreased functional connectivity with several other sites in frontal , temporal a nd occipital cortex . The most prominent of these results was decreased connectivi ty between left frontal regions classically implicated in language processing ( le ft IFG and DLPFC ) and right hemisphere regions —in the superior temporal gyrus an d sulcus , Heschl's gyrus , and anterior insula—that have been implicated in pitch processing ( Lee et al . , 2011; Garcea et al . , 2017; Warren et al . , 2003; Hoh mann et al . , 2018 ) . We suggest that this decreased connectivity between right hemisphere pitch and left hemisphere frontal cortices may relate to the unreliabil ity of the amusics' perception of and memory for pitch . This is similar to the 'w eighted connections' model of multisensory integration , where a more ( or less ) reliable modality is given a stronger ( or weaker ) weight ( Beauchamp et al . , 2 010 ) . Congenital amusia is often described as a disorder related to structural a nd functional connectivity within the right hemisphere , particularly between righ t inferior frontal cortices and right posterior temporal cortex ( see Peretz , 201 6 for review ) . Consistent with this proposal , we found in the present study tha t right inferior frontal cortex exhibited strongly decreased functional connectivi ty in the amusia group , and follow-up seed testing revealed that right auditory a reas were involved as well . However , we also found that sites in left frontal co rtex also showed large decreases in connectivity in amusia , also most prominently with right hemisphere auditory areas . Our results are consistent with an account that right hemisphere auditory areas are not only abnormally connected to right fr ontal areas ( as observed during tonal tasks ) but are less integrated with fronta 1 left hemisphere regions when processing speech and language . Our null results f or group differences in activation during speech processing are consistent with pr ior reports that amusics and controls do not differ in pitch representations withi n sensory regions . For example , the extent of pitch-responsive regions within au ditory cortex has been shown to be similar in participants with amusia and control s ( Norman-Haignere et al . , 2016 ) . Brainstem encoding of pitch in speech and m usical stimuli is similarly unimpaired in individuals with amusia ( Liu et al . , 2015b ) . Moreover , in oddball EEG paradigms , amusics show similar pre-attentive mismatch negativity responses to small pitch deviants , but impaired attention-dep endent P300 responses ( Moreau et al . , 2009; Peretz et al . , 2009; Mignault Gou let et al . , 2012; Moreau et al . , 2013 ) . These findings , along with the fact that amusics show intact non-volitional behavioral responses ( unconscious pitch s hifts ) when presented with pitch-altered feedback of their own voice ( Hutchins a nd Peretz , 2012 ) , have been interpreted as evidence that amusia is a disorder o f pitch awareness rather than one of low-level pitch processing ( Peretz et al . , 2009 ) , with differences in structural connectivity as one possible foundation of this putative impaired pitch awareness ( Hyde et al . , 2006; Loui et al . , 2009; but see Chen et al . , 2015 ) . Our interpretation of differences in functional co nnectivity between amusics and controls diverges somewhat from these previous appr oaches: we argue that down-weighting of pitch information during perceptual catego rization in both speech and music is adaptive , inasmuch as amusics have learned  $\mathsf{t}$ hat pitch is an unreliable source of evidence relative to other perceptual dimensi ons . The evidence above suggesting that encoding of pitch in the brainstem and au ditory cortex and pre-attentive responses to pitch changes are unaffected in amusi a can be interpreted as suggesting that the fundamental deficit in amusia may not be increased perceptual noise or decreased pitch awareness but difficulties with r etention of pitch information in memory ( see Tillmann et al . , 2016 for review ) . Our task arguably taxed working memory resources: in a similar paradigm performe d by the same participants in quiet listening conditions ( Jasmin et al . , 2020a ), the mean reaction time measured from the end of the second auditory stimulus w as 1 . 64 s , indicating that participants needed some time to compare both audito ry presentations and make their judgments . This interpretation is consistent with evidence suggesting that amusics have difficulty with pitch sequence processing ta sks even when discrimination thresholds are accounted for ( Tillmann et al . , 200 9 ) , as well as the finding that delaying the time interval between standard and comparison tones exacerbates pitch discrimination impairment in individuals with a musia ( Williamson et al . , 2010 ) . Moreover , the pitch awareness account of am usia cannot explain the Jasmin et al . , 2020a finding that pitch cues are downwei ghted only during longer-scale suprasegmental speech perception , while pitch weig hting is not different between amusics and controls during shorter-scale segmental speech perception , despite pitch cues being arguably more subtle in the segmental condition . However , this finding can be explained by the pitch memory account , as the suprasegmental task requires detection of and memory for pitch patterns wit hin a complex sequence , while the segmental task does not . Furthermore , an acco unt of amusia which suggests that the disorder primarily stems from differences in structural connectivity cannot account for the recent finding that functional conn ectivity patterns do not differ between amusics and controls during a verbal memor y task ( Albouy et al . , 2019 ) , as well as our finding that amusics and control s show similar functional connectivity patterns during passive listening to tone s equences . We suggest , therefore , that amusics neglect pitch because they have i mplicitly learned that their memory for pitch is unreliable , and that this down-w eighting of pitch is reflected in decreased functional connectivity between right auditory areas and downstream task-relevant areas which integrate information from

perceptual regions . One way to test this hypothesis would be to examine functiona l connectivity during perceptual categorization of consonant-vowel syllables as vo iced versus unvoiced based on a pitch cue ( F0 of the following vowel ) and a dura tional cue (voice onset time ) . We predict , based on our previous findings ( Ja smin et al . , 2020a ) , that functional connectivity will not differ between amus ics and controls on this task , a finding which would not be predicted by the pitc h awareness account of amusia . We note that a previous fMRI study on amusia detec ted group differences in functional connectivity during passive listening to tones . That study used task-defined seed voxels in bilateral auditory cortex and found , in the amusia group , increased connectivity between left and right auditory cor tex , but decreased connectivity between right auditory cortex and right inferior frontal gyrus ( Hyde et al . , 2011 ) . The present study does not necessarily cla sh with these findings , as we used different seed ROIs selected with a different procedure . We did not observe any differences in functional connectivity between conditions in our speech task . This may be because our functional imaging protoco l was timed to capture the peak in the BOLD signal corresponding to the presentati on of the second auditory stimulus . Participants never knew ( even implicitly ) w hich acoustic dimension might be useful on any given trial until after they had he ard both spoken sentences and needed to compare them to make their response . Furt hermore , pitch fluctuations in the stimuli were above participants' thresholds , even in the Duration-Informative condition ( where the standard deviation of F0 ov er each spoken utterance was , on average , 2 . 7 semitones ) , and so it is unsur prising that functional connectivity did not change on a trial-by-trial basis , an d instead the same 'neural strategy' was employed to process speech regardless of the trial type . Several other future directions are suggested by our results , pa rticularly for examining cue weighting during auditory/speech perception . In the multimodal integration studies mentioned above ( Beauchamp et al . , 2010; Nath an d Beauchamp , 2011 ) , reliability of two different sensory modalities was manipul ated experimentally by severely degrading input channels with noise , resulting in changes in connectivity . Similarly , aspects of speech could be selectively maske d with noise in order to make them less reliable , which in turn could cause corre sponding changes in functional or effective connectivity . Indeed , behavioral wor k has indicated that when fundamental frequency ( pitch ) or durational aspects of speech are manipulated to be unreliable cues , categorization behavior shifts such that participants place less relative weight on the dimension that has been made l ess reliable ( Holt and Lotto , 2010 ) . Certain groups , such as tone language sp eakers , are known to have fine-grained pitch perception abilities , and tend to p lace greater weight on pitch even when processing speech from a second , non-tonal language that they have learned ( e . g . English; Yu and Andruski , 2010; Zhang a nd Francis , 2010 , Zhang et al . , 2008; Qin et al . , 2017; Jasmin et al . , 202 0a ) . Given the increased reliability of their pitch perception , tone language s peakers may exhibit correspondingly high connectivity strength between right hemis phere auditory regions and left hemisphere 'language regions' when pitch cues are present ( more so than native non-tonal language speakers ) . Expert musicians als o have extensive pitch-related experience and training and could also serve as a p opulation to examine in future work . \n Participants , 15 individuals with amusia ( 10 F , age\xa0=\xa060 . 2\xa0±\xa09 . 4 , range\xa0=\xa043-74 ) and 15 controls ( 10 F , age\xa0=\xa061 . 3\xa0±\xa010 . 4 , range\xa0=\xa038-74 ) , were recruite d from the UK and were native British English speakers . The amusic group sample s ize reflected the maximum number of participants that could be screened and tested during our data collection period . The control group sample size was matched to t his . All participants gave informed consent , and ethical approval was obtained f rom the relevant UCL and Birkbeck ethics committees . Amusia status was obtained u sing the Montreal Battery for the Evaluation of Amusia ( MBEA ) . Participants wit h a composite score ( summing the Scale , Contour and Interval tests scores ) of 6 5 or less were classified as having amusia ( Peretz et al . , 2003 ) . We also not

e that the amusics defined using the MBEA had higher pitch thresholds than control s ( Wilcoxon Rank Sum W\xa0=\xa029 , p=0 . 001 ) but did not differ from controls in tone duration discrimination ( W\xa0=\xa0129 , p=0 . 74 ) , speech-in-noise thr eshold (  $W\times a0=\times a0155$  . 5 , p=0 . 17 ) , or audiometric hearing thresholds ( t ( 28 ) = 1 . 33 , p=0 . 20; see Jasmin et al . , 2020a for detailed methods for thes e procedures ) . The stimuli were 42 compound sentences that consisted of a pre-po sed subordinate clause followed by a main clause ( see Figure 1 for an example , a nd Jasmin et al . , 2020a , Jasmin et al . , 2020b for details ) . There were two versions of each sentence: (1) an 'early closure' version, where the verb of th e subordinate clause was used intransitively and the following noun was the subjec t of a new clause ['After Jane dusts , the dining table [is clean]"]; and ( 2 ) , 'late closure' , where the verb was transitive and took the following noun as its object , moving the phrase boundary to a slightly later position in the sentence ['After Jane dusts the dining table , [it is clean]"] . The words in both versions of the sentence were identical from the start of the sentence until the end of the second noun ( 'After Jane dusts the dining table ..." ) , and only the lexically ide ntical portions of the sentences were presented to participants; thus the two stim uli did not differ in words spoken . A native British English speaking male ( who had previously trained as an actor ) recorded early closure and late closure versi ons of each sentence in a sound-proofed room . The recordings were cropped such th at only the portions with the same words remained , and silent pauses after phrase breaks were removed . Synthesized versions of these sentences were created with ST RAIGHT voice-morphing software ( Kawahara and Irino , 2005 ) . First , the two ver sions of the sentence were manually time-aligned by marking corresponding 'anchor points' in the two recordings . Then , morphed speech was synthesized by varying t he degree to which the early closure and late closure recordings contributed durat ion and pitch information . We synthesized pairs of stimuli in three conditions: ( 1 ) In the Pitch-Informative condition , the stimulus pair had exactly the same du rational properties ( that is , the length of phonemes , syllables , and words was the average between the two original recordings ) but the vocal pitch indicated ea rly or late closure at a morphing level of 80%; ( 2 ) in the Duration-Informative condition , vocal pitch in the stimulus pair was identical ( at 50% between both v ersions ) but the durational characteristics indicated early or late closure at a morphing level of 80%; ( 3 ) in the Both-Informative condition , both pitch and ti me cued early or late closure simultaneously at 80% . The morphed speech varied on ly in duration and pitch , while all other aspects of the acoustics ( such as ampl itude and spectral characteristics other than pitch ) were the same , held constan t at 50% between the two original recordings during morphing . This stimulus set i s freely available ( Jasmin et al . , 2020b ) . Across all stimuli , F0 ( vocal pi tch ) differences between early and late closure versions were large , with a mean of maximum difference of 7 . 7 semitones and range of 4 . 0-12 . 6 semitones . Thu s , even the stimulus pair with the smallest pitch difference ( 4 . 0 semitones ) exceeded the \xa0~1 . 5 semitone pitch change detection threshold of the 'most impa ired' participant in the amusia group ( Jasmin et al . , 2020a ) , which increased the chances that the amusia group would not suffer from poor performance , thereby avoiding a performance-related confound with our experimental design ( see Church et al . , 2010 for discussion ) . Subjects were scanned with a Siemens Avanto 1 . 5 Tesla magnetic resonance imaging scanner with a 32-channel head coil , with soun ds presented via Sensimetrics S14 earbuds , padded around the ear with NoMoCo memo ry foam cushions . Functional data were collected using a slow event-related desig n with sparse temporal sampling to allow presentation of auditory stimuli in quiet . We used an echo planar image sequence , with 40 slices , slice time 85 ms , slab tilted to capture the entire cerebrum and dorsal cerebellum , ascending sequential acquisition; 3\xa0x\xa03\xa0x\xa03 mm voxel size; silent stimulus and response per iod\xa0=\xa08 . 7 s , volume acquisition time\xa0=\xa03 . 4 s , total inter-trial interval\xa0=\xa012 . 1 s , flip angle\xa0=\xa090 degrees , bandwidth\xa0=\xa02298

Hz/pixel , echo time ( TE ) =\xa050 ms . After collecting functional runs , a high -resolution T1-weighted structural scan was collected ( MPRAGE , 176 slices , sagi ttal acquisition , 2x GRAPPA acceleration , 1 mm isotropic voxels , acquisition ma trix\xa0=\xa0224\xa0x\xa0256 ) . Each run began with three dummy scans to allow ma gnetic stabilization . Each trial ( repetition time ) lasted 12 . 1 s . The start of each trial was triggered by a pulse corresponding to the start of a volume acqu isition ( which acquired neural data from the previous trial , at a delay ) . At t \xa0=\xa01 s into the trial , the sentence appeared on the screen; before scanning participants were instructed to read each sentence silently to themselves . At t\x a0=\xa05 s ( plus or minus a random 100 ms jitter ) participants heard a spoken ve rsion of the first part of the sentence . At  $t\a0=\x00$  . 4 s ( plus or minus 100  $\mbox{\it ms}$  jitter ) the second version was presented . The two spoken versions contained t he same words but their pitch and/or timing characteristics cued a phrase boundary that occurred earlier or later in the sentence . Following this , there were appro ximately 2 s of silence during which the participant responded with the button box , before the scanner began acquiring the next volume at t\xa0=\xa012 . 1 s . Parti cipants performed three blocks of 42 trials ( 14 each of Pitch-Informative , Durat ion-Informative , and Both-Informative ) with 8 Rest trials interspersed within ea ch block . Following data collection for this task and the structural scan , parti cipants took part in two task-free fMRI scanning runs in which they watched a sile nt film ( The General , starring Buster Keaton , or an episode of the Planet Earth series played without sound ) while being presented auditorily with semi-random to ne sequences . Stimuli consisted of sequences of 'pips' - 30 ms 6-harmonic complex tones . The fundamental frequencies of the pips were either 440 , 466 . 16 , 493 . 88 or 523 . 25 Hz , and the time between tone onsets was 0 . 075 , 0 . 125 , 0 . 175 , or 0 . 225 s . The transition probabilities ( determining whether pip N+1 had the same pitch or duration properties as pitch N ) were set at either 0 . 1 and 0 . 9 for duration and either 0 . 3 and 0 . 7 for pitch . These two transition param eters were 'crossed' to create four design cells , and 25 random sequences were ge nerated for each cell . MRI scanning parameters were identical to those used in th e active , prosody task , except the time between volume acquisitions was 17 . 1 s . Participants listened to 100 tone sequences across two runs ( 50 per run ) . Mat lab code used to create the stimuli can be found online ( see Data Availability St atement ) . Image preprocessing was performed with FreeSurfer 6 . 0 . 0 (Fischl, 2012 ) and AFNI-SUMA 18 . 1 . 18 ( Cox , 1996 ) . Anatomical images were registere d to the third echo planar image of the first run using Freesurfer's bbregister an d processed with FreeSurfer's automated pipeline for segmenting tissue types , gen erating cortical surface models , and parcellating subcortical structures . Masks of inferior colliculi were obtained by manually examining individual subjects' ana tomical images and selecting a single EPI voxel located at its centre , bilaterall y . Freesurfer cortical surface models were imported to AFNI with the @SUMA\_Make\_S pec\_FS program . Then a standard pre-processing pipeline using AFNI's afniproc . p y program was used: all echo planar image volumes were aligned to the third repeti tion time of the first run using AFNI's 3DAllineate , intersected with the cortica 1 surface with SUMA , smoothed along the surface with a 2D 6-mm-FWHM kernel , and converted to a standard mesh ( std . 141 ) for group analyses , separately for eac h hemisphere , where each vertex in the mesh ( 198812 per hemisphere ) is aligned to the 'same' location in the cortex across subjects , using curvature-based morph ing . Preprocessing of the passive listening experiment data was identical . The m agnitude of transient head motion was calculated from the six motion parameters ob tained during image realignment and aggregated as a single variable using AFNI's @ 1dDiffMag to calculate a Motion Index ( Berman et al . , 2016; Gotts et al . , 201 2; Jasmin et al . , 2019 ) . This measure is similar to average Frame Displacement over a scan ( Power et al . , 2012 ) and is in units of mm per repetition time . T he difference in average motion between the groups was small ( amusia group mean m otion\xa0=\xa00 . 31 mm/TR; control group mean\xa0=\xa00 . 28 mm/TR ) and amounted

to 32 micrometers (  $\sim$ 1/30th of a millimeter ) per TR . The mean and distribution o f motion did not differ statistically between groups ( two sample t-test p=0 . 70 , two-tailed ) . Given the previous reports ( described above ) of changes in conn ection strength between unimodal and multimodal areas in response to noise ( Beauc hamp et al . , 2010; Nath and Beauchamp , 2011 ) , we chose a connectivity-based a nalysis approach for our study . Beta series correlation ( Rissman et al . , 2004 ) is a technique for examining functional connectivity and its modulation by task , using correlations in trial-by-trial responses . It has been shown to be more po werful than alternatives such as generalized psycho-physiological interaction ( gP PI ) for event-related designs ( Cisler et al . , 2014 ) . In a beta series analys is , one beta weight is calculated for each trial in the experiment ( rather than for each condition ) . All the trial-wise betas associated with a given condition are then serially ordered to form a 'beta series' . Finally , using the beta serie s in the same way as a standard BOLD fMRI time series , functional connectivity ( measured as Pearson correlations ) is calculated between seed regions of interest and the rest of the brain . Differences in functional connectivity can then be exa mined by comparing groups , comparing conditions , or examining the interaction of these factors . Our experiment used a slow event-related design with a long repeti tion time ( 12 . 1 s ) and sparse temporal sampling ( with volume acquisition sepa rated by silent periods ) . Therefore , the time between acquisitions was long eno ugh for the haemodynamic response to return to baseline , and each echo planar ima ge acquisition corresponded to exactly one trial ( Figure 1 ) . For this reason , we did not convolve the echo planar image time series with a basis function during subject-level statistical analysis ( Hall et al . , 1999 ) . In the design matrix for obtaining trial-wise betas , 126 column regressors were used ( one for each no n-rest trial ) . Each column vector was of length 150 ( corresponding to all trial s , including rest trials ) and had a single 'one' in the position where the trial associated with that column occurred , while zeros were located in every other pos ition . Polynomials up to second degree were also included in the model , on a run -wise basis , to remove the mean and any linear or quadratic trends . Fitting the trial regressors on a subject-wise basis resulted in cortical surface models of be ta weights for each of the 126 trials , at each vertex on the reduced-vertex icosa hedral cortical surface , with beta weights reflecting the neural response associa ted with that trial . As noted above , trial-wise betas were then serially ordered to form beta series separately for each of the three experimental conditions ( Pit ch-Informative , Duration-Informative , and Both-Informative ) ( Rissman et al . , 2004 ) . Because there were 30 participants , this procedure resulted in a total o f 90 beta series ( 30 participants\xa0x\xa03 conditions=90 beta series ) . As for the passive tone listening data , because all 'trials' were of the same type , it was not necessary to separate them into conditions and perform a first-level model to obtain betas . However , polynomials up to second degree were detrended from th e pre-processed data ( as was done with the task data ) . Beta series analysis req uires initial seed voxels , vertices , or regions to be identified , whose trial-t o-trial changes in activity are then compared to those of the rest of the brain . Rather than choose a priori seeds derived from the literature , which used mainly musical tasks or resting state , we used a data-driven approach to search for the largest group and condition differences in functional connectivity ( Berman et al ., 2016; Cole et al ., 2010; Gotts et al ., 2012; Jasmin et al ., 2019; Meoded et al . , 2015; Song et al . , 2015; Steel et al . , 2016; Stoddard et al . , 201 6; Watsky et al . , 2018 ) . To do this , we first calculated the 'whole-brain con nectedness' of each cortical vertex ( a procedure available in AFNI as the 3dTCorr Map function ) . The whole-brain connectedness of a given vertex is defined as the Pearson correlation of activity within that vertex/voxel and the average signal ac ross all neural gray matter in the rest of the brain . Mathematically , this is eq uivalent to calculating thousands of Pearson correlations , of a given vertex/voxe 1 series and every other vertex/voxel series in the brain , and then taking the me

an of those correlations ( Cole et al . , 2010 ) , then repeating the process for every individual voxel/vertex . As such , it represents the global connectedness ( or 'global correlation' ) of a vertex/voxel . To calculate whole-brain connectedne ss , first , the average of trial-wise betas in gray matter across the brain was c alculated in volume space , separately for each subject and for each condition , b y running first-level ( subject ) models . The statistical models were identical t o those conducted on the cortical surface , described above , but were performed o n volumetric Talairach images instead of the cortical surfaces . The reason for th is choice was so that voxels in cortex and subcortex would contribute equally to o ur measure of global ( whole-brain ) connectivity . First , the\xa0average gray-ma tter beta value was calculated for each trial by intersecting each image in the be ta series with a whole-brain gray matter mask ( which excluded white matter and ve ntricles ) and calculating the average beta value within the mask ( Gotts et al . , 2012; Jasmin et al . , 2019 ) . Next , this gray matter average was correlated w ith each cortical surface vertex's beta series , separately for each subject and c ondition , to obtain whole-brain connectedness maps . These values were then subje cted to a statistical analysis based on our 2 ( Group ) ×\xa03 ( Condition ) exper imental design . Linear mixed effects models ( AFNI's 3dLME ) ( Chen et al . , 201 3 ) were constructed whose dependent variables were the vertex-wise whole-brain co nnectedness maps from each beta series . Group and Condition and their interaction were included as fixed effects . Participant was treated as a random intercept . R esults of this step were corrected vertex-wise for multiple comparisons with False Discovery Rate ( q\xa0<\xa00 . 05 ) , separately for each test ( Main Effect of Gr oup; Main Effect of Condition; Interaction of Group by Condition ) by pooling the p-values from both hemispheres' cortical surfaces . This False Discovery Rate thre shold corresponded to uncorrected p<4 $\times$ a0 $\times$ a010-6 for the Main Effect of Group . Four significant results (contiguous significant vertices) survived this thresho ld and were taken forward for the next analysis step . For the Main Effect of Cond ition and Interaction of Condition x Group , no results survived statistical corre ction at  $\advantum{1}{xa0FDR}$  (  $\advantum{1}{q}\advantum{2}{xa00}$  . 05 ) . An analogous procedure was run on the passi ve tone listening data , in which whole-brain connectedness values were compared b y group ( amusic vs . control ) in a linear mixed effects model . No significant F DR-corrected group differences were detected , nor at a reasonable uncorrected thr eshold of p<0 . 001 . A similar procedure was performed for subcortical structures . Beta series were obtained for each subject , structure , and experimental condit ion , from their standard Freesurfer subcortical parcellations by masking the EPI data within each structure and calculating the average of the voxels . Each struct ure's beta series was then correlated with the whole-brain gray matter beta averag e , separately for each condition , and the resulting values were subjected to lin ear mixed effects models with the same factors as above . Tests for Main Effect of Condition , of Group , and the Interaction of these factors was performed . All pvalues were greater than p>0 . 001 and no results survived an FDR-correction calcu lated over them . The first analysis step ( seed definition , described above ) id entified which , if any , brain areas showed the largest connectivity differences between groups . However , this step is insufficient to localize the other specifi c regions driving this pattern . An analogy is in Analysis of Variance , where a s ignificant omnibus test indicates a difference exists , but follow-up testing is r equired to determine where in the model differences exist ( Gotts et al . , 2012 ) . Thus , to locate the regions driving this pattern , we undertook a second step: follow-up seed-to-whole-brain testing ( Cole et al . , 2010; Gotts et al . , 2012; Jasmin et al . , 2019 ) . Each seed region was examined with respect to its connec tivity pattern with every cortical vertex and subcortical structure . For each of the 90 beta series ( 30 subjects by three conditions ) , values within the seed ve rtices were averaged and then correlated with the beta series for every vertex in the brain . These correlations were Fisher Z-transformed and used as the dependent variables in linear mixed effects models ( 3dLME ) with the same fixed and random

effects as above . For each of the seeds , we tested for the group difference ( Am usia vs Control ) in connectivity . Results were False Discovery Rate corrected to ( q\xa0<\xa00 . 05 ) across all eight follow-up tests [four seeds\xa0×\xa02 hemisp heres] corresponding to a threshold of p<0 . 00035 . Similarly , for the subcortic al structures , each seed beta series was correlated with subcortical structure be ta series , with resulting values subject to statistical testing . An FDR correcti on over all tests involving subcortex was applied . For display in figures , the d ata were converted from SUMA's standard mesh ( std . 141 ) to Freesurfer's standar d surface (fsaverage) using AFNI's SurfToSurf program and mapping values from th e closest nodes ( i . e . vertices ) . To determine whether the functional connect ivity patterns we observed were related to the importance placed on acoustic dimen sions during prosodic categorization ( cue weighting ) , the functional connectivi ty results were analyzed with respect to previously acquired cue weights obtained behaviorally from a subset of participants ( Jasmin et al . , 2020a ) . The right anterior insula and right auditory cortex results were used as ROIs (Figure 3A) . The beta series for each ROI ( averaged across vertices ) was correlated with th e beta series within the L-DLPFC seed area , separately for each condition , then averaged and Fisher Z-transformed . For the 21 participants for whom we had prosod ic cue weight data ( from Jasmin et al . , 2020a ) , these cue weights were analyz ed with respect to the functional connectivity between the L-DLPFC seed and the tw o ROIs using Spearman correlations . As described above , functional connectivity between L-DLPFC , and right auditory cortex and right insula was calculated using data from the passive tone listening task , using ROIs derived from the active spe ech perception task . After pre-processing and de-trending , the averaged value fr om the tone listening experiment within these ROIs was extracted , as well as the LDLPFC seed , for each experiment . Correlations between signal within the seed an d the two ROIs was calculated and Fisher Z-transformed . As mentioned above , beca use all trials in the tone-listening experiment were analyzed as the same type , i t was not necessary to use a first-level model to obtain trial-wise betas . Simila rly for the data from the speech task , the average value within the seed region a nd both ROIs was extracted , separately for each of the 3 Beta series ( Pitch- , T ime- and Both-Informative ) , and the seed and ROI series were correlated . The  $\ensuremath{\mathsf{me}}$ an of these three correlation coefficients was calculated and Fisher Z-transformed . Finally , statistics were performed using a mixed ANOVA with Experiment ( Speech or Tones ) as the within-subject factor and Group ( Amusia or Control ) as the bet ween-subject factor . A standard General Linear Model comparing activation strengt h ( rather than connectivity ) was also conducted . As in the General Linear Model for obtaining beta weights , no basis function was used , and polynomials up to se cond degree were included in the models . The data that support the findings of th is study are openly available in the Birkbeck repository ( https://researchdata . bbk . ac . uk/65/ ) , as are the speech stimuli ( Jasmin et al . , 2020b; https:// researchdata  $\cdot$  bbk  $\cdot$  ac  $\cdot$  uk/37/ )  $\cdot$  The speech task can be demoed at the followin g link: ( Gorilla Open Materials; https://gorilla . sc/openmaterials/102786 ) ."

```
In [15]: # test summarizing the abstract
paras = item.article.split("\n")
abstract = paras[0]
abstract
```

Out[15]: 'Individuals with congenital amusia have a lifelong history of unreliable pitch processing. Accordingly, they downweight pitch cues during speech perception and instead rely on other dimensions such as duration. We investigated the neural basis for this strategy. During fMRI, individuals with amusia (N = 15) and controls (N = 15) read sentences where a comma indicated a grammatical phrase boundary. They then heard two sentences spoken that differed only in pitch and/or duration cues and selected the best match for the written sentence. Prominent reductions in functional connectivity were detected in the amusia group between left prefrontal language-related regions and right hemisphere pitch-related regions, which reflected the between-group differences in cue weights in the same groups of listeners. Connectivity differences between these regions were not present during a control task. Our results indicate that the reliability of perceptual dimensions is linked with functional connectivity between frontal and perceptual regions and suggest a compensatory mechanism.'

```
In [16]: print("Summary based on abstract:\n----")
send_sumarize_request(abstract)

Summary based on abstract:
```

Pausing for 10 secs...

Out[16]: 'Individuals with congenital amusia, a lifelong condition characterized by unrelia ble pitch processing, rely on other dimensions such as duration during speech perc eption. A study investigating the neural basis for this strategy used fMRI to comp are functional connectivity in individuals with amusia (N = 15) and controls (N = 15) as they read sentences with grammatical phrase boundaries and then heard two s entences that differed only in pitch and/or duration cues, selecting the best match for the written sentence. Results showed prominent reductions in functional connectivity in the amusia group between left prefrontal language-related regions and right hemisphere pitch-related regions, reflecting the between-group differences in cue weights. These connectivity differences were not present during a control task, indicating a specific compensation mechanism. The study suggests that the reliability of perceptual dimensions is linked with functional connectivity between frontal and perceptual regions and that individuals with amusia develop a compensatory mechanism to rely on other dimensions such as duration during speech perception due to their unreliable pitch processing.'

```
In [17]: print("Summary based on full article:\n-----")
    s_full = send_sumarize_request(item.article)
    print(len(s_full))
    print(len(s_full.split()))
    print(s_full)
```

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Summary based on full article:
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Pausing for 10 secs...

OK

4484

633

Congenital amusia is a rare condition characterized by a lifelong history of unrelia ble pitch processing, resulting in a strategy of downweighting pitch cues during spe ech perception and relying on other dimensions such as duration. A study using fMRI found that individuals with amusia exhibited prominent reductions in functional conn ectivity between left prefrontal language-related regions and right hemisphere pitch -related regions, reflecting the between-group differences in cue weights. These con nectivity differences were not present during a control task. The results suggest a compensatory mechanism for the reduced reliability of perceptual dimensions. While c ongenital amusia is believed to be innate, recovery is possible through training. Pi tch is important for cueing categories in spoken language, conveying emotion in spee ch, and is usually associated with music. In highly controlled laboratory tasks, onl y minor deficits have been observed in amusia, and in naturalistic speech perception contexts, people with amusia rarely report any difficulties. This may be because, in natural speech, pitch variation tends to co-occur with variation in other acoustic d imensions. Our lab has shown that in such cases where multiple redundant cues are av ailable, English-speaking individuals with amusia tend to rely less on pitch than no n-amusic controls, suggesting they may calibrate their perception by down-weighting the cues that are less reliable for them. As for emotional prosody in speech, indivi duals with amusia can recognize emotions in spoken sentences but not in short sample s such as isolated vowels. It is unknown how decreased reliance on a particular acou stic cue during speech perception is reflected in the brain. Previous neural studies of cue integration have focused on integration of multiple modalities and the 'weigh ted connections' model of multisensory integration, where the relative reliability o f the modalities involved with perception of a stimulus is related to differential c onnectivity strength. For example, when participants simultaneously view and feel to uches to the hand, and reliability of visual and tactile perception is manipulated e xperimentally via the introduction of noise, connection strength between unimodal an d multimodal sensory areas adjusts accordingly. Similarly, effective connectivity be tween the superior temporal sulcus and visual and auditory areas has shown modulatio ns during processing of audiovisual speech, where connection strength between audito ry cortex and the STS is weaker when noise has been introduced to the auditory speec h, and conversely connection strength between visual cortex and STS is weaker if vis ual noise is introduced. An analogous phenomenon may be at work within a single moda lity during multidimensional integration, such as the acoustic speech signal carryin g multiple co-occurring acoustic dimensions which often provide redundant cues to di sambiguate a linguistic category. Individuals with typical pitch perception have lea rned through a lifetime of experience with speech acoustics that vocal pitch is a us eful and reliable cue, but individuals with amusia, who have unreliable perception o f and memory for pitch, would have learned that, for them, pitch is not a reliable c ue for processing spoken language. Thus, by analogy to the multisensory weighting re sults, it is hypothesized that amusics may exhibit decreased connectivity between la nguage regions and pitch-related areas during speech processing. The neural foundati ons of perceptual weighting in speech have not been investigated in atypical individ uals. A study using fMRI scanned 15 individuals with amusia and 15 controls and foun d that during speech encoding, there were no group differences in task-related activ ation or functional connectivity during processing of speech. However, functional co nnectivity between these areas was analyzed with respect to prosodic cue weights obt ained outside the scanner and showed that across this subset of participants, normal ized cue weights were correlated with L-DLPFC<=>R insula connectivity and L-DLPFC<=> R auditory cortex connectivity, indicating that participants who relied least on pit

ch information to process speech had the weakest functional connectivity between the se areas, while those who relied most on pitch had the strongest. These results sugg est that even non-amusics may perform dimensional reweighting of acoustic dimensions and functional connectivity.

Out[19]: 'Spoken language is colored by fluctuations in pitch and rhythm . Rather than spea king in a flat monotone , we allow our sentences to rise and fall . We vary the le ngth of syllables , drawing out some , and shortening others . These fluctuations , known as prosody , add emotion to speech and denote punctuation . In written lan guage , we use a comma or a period to signal a boundary between phrases . In speec h , we use changes in pitch - how deep or sharp a voice sounds - or in the length of syllables . Having more than one type of cue that can signal emotion or transit ions between sentences has a number of advantages . It means that people can under stand each other even when factors such as background noise obscure one set of cue s . It also means that people with impaired sound perception can still understand speech . Those with a condition called congenital amusia , for example , struggle to perceive pitch , but they can compensate for this difficulty by placing greater emphasis on other aspects of speech . Jasmin et al . showed how the brain achieves this by comparing the brain activities of people with and without amusia . Partici pants were asked to read sentences on a screen where a comma indicated a boundary between two phrases . They then heard two spoken sentences , and had to choose the one that matched the written sentence . The spoken sentences used changes in pitch and/or syllable duration to signal the position of the comma . This provided liste ners with the information needed to distinguish between "after John runs the race , . . . " and "after John runs , the race . . . " , for example . When two brain  ${\sf r}$ egions communicate , they tend to increase their activity at around the same time . The brain regions are then said to show functional connectivity . Jasmin et al . found that compared to healthy volunteers , people with amusia showed less functio nal connectivity between left hemisphere brain regions that process language and r ight hemisphere regions that process pitch . In other words , because pitch is a l ess reliable source of information for people with amusia , they recruit pitch-rel ated brain regions less when processing speech . These results add to our understa nding of how brains compensate for impaired perception . This may be useful for un derstanding the neural basis of compensation in other clinical conditions . It cou ld also help us design bespoke hearing aids or other communication devices , such as computer programs that convert text into speech . Such programs could tailor th e pitch and rhythm characteristics of the speech they produce to suit the percepti on of individual users .'

```
In [20]: # apply to all rows in eval miniset
    text_cap = 20_000 # temporarily limit to 20k characters, set to -1 for full text
    print("Summarization process started...")
```

df["groq\_mistral\_summary"] = df["article"].apply(lambda text: send\_sumarize\_request
print("Completed")

Summarization process started... Sending request for text = Mature neural networks synchronize and integrate spatiote mporal activity patterns to support cogniti Pausing for 10 secs... OK Sending request for text = Many decisions are thought to arise via the accumulation of noisy evidence to a threshold or bound . Pausing for 10 secs... OK Sending request for text = Mutations in the inositol 5-phosphatase OCRL cause Lowe s yndrome and Dent's disease . Although OCRL Pausing for 10 secs... OK Sending request for text = Gradients of signaling proteins are essential for inducin g tissue morphogenesis . However , mechanis Pausing for 10 secs... OK Sending request for text = Similarity between two individuals in the combination of genetic markers along their chromosomes ind Pausing for 10 secs... OK Sending request for text = Latent Epstein-Barr virus ( EBV ) infection is causally 1 inked to several human cancers . EBV expres Pausing for 10 secs... Sending request for text = Dynamic post-translational modification of RNA polymerase II ( RNAPII ) coordinates the co-transcrip Pausing for 10 secs... Sending request for text = Swi2/Snf2 ATPases remodel substrates such as nucleosomes and transcription complexes to control a wi Pausing for 10 secs... OK Sending request for text = Accurate chromosome segregation depends on coordination b etween cohesion resolution and kinetochore-Pausing for 10 secs... Sending request for text = Streptococcus pneumoniae is a leading cause of invasive d isease in infants , especially in low-incom Pausing for 10 secs... OK Sending request for text = C4 photosynthesis has independently evolved from the ance stral C3 pathway in at least 60 plant linea Pausing for 10 secs... Sending request for text = Host shutoff is a common strategy used by viruses to repr ess cellular mRNA translation and concomita Pausing for 10 secs... Sending request for text = The transcription factor RpaA is the master regulator of circadian transcription in cyanobacteria, Pausing for 10 secs... OK Sending request for text = Coiled coils are the best-understood protein fold , as th

eir backbone structure can uniquely be desc

```
Sending request for text = Hemoglobin ( Hb ) represents a model protein to study mol
        ecular adaptation in vertebrates . Although
        Pausing for 10 secs...
        OK
        Sending request for text = We report a functional switching valve within the female
        genitalia of the Brazilian cave insect Neot
        Pausing for 10 secs...
        Sending request for text = Previous studies tracking AMPA receptor ( AMPAR ) diffusi
        on at synapses observed a large mobile extr
        Pausing for 10 secs...
        OK
        Sending request for text = Histone acetylation and deposition of H2A . Z variant are
        integral aspects of active transcription .
        Pausing for 10 secs...
        OK
        Sending request for text = Membrane nanodomains have been implicated in Ras signalin
        g , but what these domains are and how they
        Pausing for 10 secs...
        Sending request for text = This is an analysis of how magnetic fields affect biologi
        cal molecules and cells . It was prompted b
        Pausing for 10 secs...
        Sending request for text = Individuals with congenital amusia have a lifelong histor
        y of unreliable pitch processing . Accordin
        Pausing for 10 secs...
        Sending request for text = NCOA4 is a selective cargo receptor for the autophagic tu
        rnover of ferritin , a process critical for
        Pausing for 10 secs...
        OK
        Sending request for text = ISG15 is an interferon-stimulated , linear di-ubiquitin-l
        ike protein , with anti-viral activity . Th
        Pausing for 10 secs...
        Sending request for text = Temporal experience of odor gradients is important in spa
        tial orientation of animals . The fruit fly
        Pausing for 10 secs...
        OK
        Completed
In [21]: # check how many rows have blank result (some errors)
         empty_df = df.query("groq_mistral_summary.str.strip() == ''")
         print(len(empty_df))
        0
In [22]: # attempt to retry for blank rows (due to some errors)
         retry = True
         if retry:
             print("Retrying for empty results...")
             for i in range(len(df)):
                 item = df.iloc[i]
```

if item["groq\_mistral\_summary"] == "":

```
print("Item =", i)
    text = item["article"]
    df.at[i, "groq_mistral_summary"] = send_sumarize_request(text[:text_cap
print("Completed")
```

Retrying for empty results...
Completed

```
In [23]: df
```

Out[23]:		lay_summary	article	headings	keywords	id	groq_mistral_summ
	0	It can take several months , or even years , f	Mature neural networks synchronize and integra	[Abstract, Introduction, Results, Discussion, 	[neuroscience]	elife- 69011- v2	The developmer neural networks f an im
	1	Many of our decisions are made on the basis of	Many decisions are thought to arise via the ac	[Abstract, Introduction, Results, Discussion, 	[neuroscience]	elife- 17688- v1	The article discu how decisions are m
	2	Oculo-Cerebro- Renal syndrome of Lowe ( Lowe sy	Mutations in the inositol 5- phosphatase OCRL c	[Abstract, Introduction, Results, Discussion, 	[cell biology]	elife- 02975- v2	Mutations in inositol 5-phospha OCRI
	3	When an embryo develops, its cells must work	Gradients of signaling proteins are essential	[Abstract, Introduction, Results, Discussion, 	[developmental biology]	elife- 38137- v3	The distributio signaling proteins,
	4	Our genomes contain a record of historical eve	Similarity between two individuals in the comb	[Abstract, Introduction, Results, Discussion, 	[evolutionary biology, genetics and genomics]	elife- 15266- v1	The article discusses use of genetic mar
	5	Over 90% of adults around the world are infect	Latent Epstein- Barr virus ( EBV ) infection is	[Abstract, Introduction, Results, Discussion, 	[microbiology and infectious disease, cancer b	elife- 22509- v2	Epstein-Barr virus (E is a widespr hum
	6	Genes are sections of DNA that encode the inst	Dynamic post- translational modification of RNA	[Abstract, Introduction, Results, Discussion, 	[chromosomes and gene expression, computationa	elife- 11215- v2	The C-terminal don (CTD) of F polymeras
	7	An organism's DNA contains thousands of genes	Swi2/Snf2 ATPases remodel substrates such as n	[Abstract, Introduction, Results, Discussion, 	[structural biology and molecular biophysics]	elife- 07432- v2	The Swi2/Snf2 ATPa are a large and divi
	8	Human reproductive cells—eggs and sperm—are pr	Accurate chromosome segregation depends on coo	[Abstract, Introduction, Results, Discussion, 	[chromosomes and gene expression, cell biology]	elife- 01133- v1	In mammalian oocy Shugoshin-like pro
	9	Microorganisms live in most	Streptococcus pneumoniae is	[Abstract, Introduction,	[microbiology and infectious	elife- 26255-	The bacter Streptoco

	lay_summary	article	headings	keywords	id	groq_mistral_summ
	parts of our body	a leading cause of	Results, Discussion, 	disease, genetics	v2	pneumoniae, a le
10	Plants rely on carbon for their growth and sur	C4 photosynthesis has independently evolved fr	[Abstract, Introduction, Results, Discussion, 	[plant biology]	elife- 00961- v1	C4 photosynthes complex trait foun
11	Proteins carry out diverse activities in our C	Host shutoff is a common strategy used by viru	[Abstract, Introduction, Results, Discussion, 	[computational and systems biology, microbiolo	elife- 18311- v2	Influenza A virus (IA' a negative-sense
12	The cycle of day and night is one of the most 	The transcription factor RpaA is the master re	[Abstract, Introduction, Results, Discussion, 	[short report, biochemistry and chemical biolo	elife- 23210- v1	The transcription fa RpaA is the master
13	Proteins are made up of building blocks called	Coiled coils are the best- understood protein f	[Abstract, Introduction, Results and discussio	[structural biology and molecular biophysics]	elife- 11861- v2	Coiled coils are v understood pro fol
14	In humans and other mammals , a protein in the	Hemoglobin ( Hb ) represents a model protein t	[Abstract, Introduction, Results, Discussion, 	[biochemistry and chemical biology]	elife- 47640- v1	Hemoglobin (Hb) tetrameric allost p
15	In dry caves of southeastern Brazil , live a g	We report a functional switching valve within	[Abstract, Introduction, Results, Discussion, 	[evolutionary biology]	elife- 39563- v2	The Brazilian cave in Neotrogla has a fu
16	Forgetting is a common experience in our every	Previous studies tracking AMPA receptor ( AMPA	[Abstract, Introduction, Results, Discussion, 	[structural biology and molecular biophysics,	elife- 27744- v3	The study used superesolution microsc
17	Cells contain a large number of proteins that 	Histone acetylation and deposition of H2A . Z	[Abstract, Introduction, Results, Discussion, 	[chromosomes and gene expression, genetics and	elife- 56325- v1	In Drosophila, Domino (D chromatin re
18	The Ras family of proteins play an important r	Membrane nanodomains have been implicated in R	[Abstract, Introduction, Results, Discussion, 	[structural biology and molecular biophysics,	elife- 46393- v2	The paper reveals I membrane organiza

		lay_summary	article	headings	keywords	id	groq_mistral_summ	
	19	How biological systems interact with magnetic	This is an analysis of how magnetic fields aff	[Abstract, Introduction, Results, Discussion, 	[short report, physics of living systems, neur	elife- 17210- v3	The analysis preser here evaluates red	
	20	Spoken language is colored by fluctuations in 	Individuals with congenital amusia have a life	[Abstract, Introduction, Results, Discussion, 	[neuroscience]	elife- 53539- v2	Individuals v congenital amus condi	
	21	The cells of nearly all organisms need iron as	NCOA4 is a selective cargo receptor for the au	[Abstract, Introduction, Results, Discussion, 	[cell biology]	elife- 10308- v1	NCOA4 is a selec cargo receptor med	
	22	Listeria monocytogenes is a bacterium that can	ISG15 is an interferon- stimulated , linear di	[Abstract, Introduction, Results, Discussion, 	[cell biology, microbiology and infectious dis	elife- 06848- v1	ISG15 is an interfer stimulated, linear di	
	23	Fruit flies are attracted to the smell of rott	Temporal experience of odor gradients is impor	[Abstract, Introduction, Results, Discussion, 	[short report, neuroscience]	elife- 06651- v2	The fruit fly, Drosop melanogaster, is c	
	<pre>output_path = "./data/output/mini_dev_set/" output_filename = "elife_groq_mistral_summary.csv"  print("Writing to file ", output_filename) df.to_csv(output_path+output_filename,</pre>							
In [25]:	<pre>output_filename = "elife_groq_mistral_summary.json"  print("Writing to file ", output_filename) df.to_json(output_path+output_filename,</pre>							
		ing to file ela leted	ife_groq_mistr	al_summary.js	son			
In [26]:	<pre># process the PLOS dataset: dev_df_filename = "//data/mini_dataset/PLOS_val_mini.jsonl"</pre>							

Out[26]:		lay_summary	article	headings	keywords	id
	0	Yersinia pestis , the bacterial agent of plagu	Fleas can transmit Yersinia pestis by two mech	[Abstract, Introduction, Results, Discussion, 	[united states, invertebrates, medicine and he	journal.ppat.1006859
	1	The genome of all vertebrates is heavily colon	Endogenous retroviruses ( ERVs ) are remnants	[Abstract, Introduction, Results, Discussion, 	[viruses, sheep, virology]	journal.ppat.0030170
	2	The molecular mechanisms underlying directed c	The Drosophila embryonic gonad is assembled fr	[Abstract, Introduction, Results, Discussion, 		journal.pgen.1003720
	3	Contrary to the long- standing belief that no n	Recently , we presented a study of adult neuro	[Abstract, Introduction, Model, Results, Discu	[computational biology/computational neuroscie	journal.pcbi.1001063
	4	Embryonic stem cells have two remarkable prope	Understanding the transcriptional regulation o	[Abstract, Introduction, Results, Discussion, 	[developmental biology, cell biology, mammals,	journal.pgen.0030145
In [27]:			rows in eval m 00 # temporar		20k characters, set	to -1 for full text
	df	•		•	].apply(lambda text:	send_sumarize_reques

```
Summarization process started...
Sending request for text = Fleas can transmit Yersinia pestis by two mechanisms , ea
rly-phase transmission ( EPT ) and biofilm-
Pausing for 10 secs...
OK
Sending request for text = Endogenous retroviruses ( ERVs ) are remnants of ancient
retroviral infections of the host germline
Pausing for 10 secs...
OK
Sending request for text = The Drosophila embryonic gonad is assembled from two dist
inct cell types , the Primordial Germ Cells
Pausing for 10 secs...
OK
Sending request for text = Recently , we presented a study of adult neurogenesis in
a simplified hippocampal memory model . The
Pausing for 10 secs...
OK
Sending request for text = Understanding the transcriptional regulation of pluripote
nt cells is of fundamental interest and wil
Pausing for 10 secs...
OK
Sending request for text = The current model of hepatitis C virus ( HCV ) production
involves the assembly of virions on or nea
Pausing for 10 secs...
Sending request for text = Secondary amphiphilicity is inherent to the secondary str
uctural elements of proteins . By forming e
Pausing for 10 secs...
Sending request for text = Herein , we studied a virulent isolate of the leading bac
terial pathogen Streptococcus pneumoniae in
Pausing for 10 secs...
OK
Sending request for text = HIV is known to spread efficiently both in a cell-free st
ate and from cell to cell , however the rel
Pausing for 10 secs...
Sending request for text = An increasing number of genetic variants have been identi
fied for many complex diseases . However ,
Pausing for 10 secs...
OK
Sending request for text = The Saccharomyces cerevisae RAD3 gene is the homolog of h
uman XPD , an essential gene encoding a DNA
Pausing for 10 secs...
Sending request for text = A budget proposal to stop the U . S . Centers for Disease
Control and Prevention ( CDC ) funding in
Pausing for 10 secs...
OK
Sending request for text = Insulator or enhancer-blocking elements are proposed to p
lay an important role in the regulation of
Pausing for 10 secs...
OK
Sending request for text = Ebolaviruses , highly lethal zoonotic pathogens , possess
longer genomes than most other non-segment
Pausing for 10 secs...
```

OK Sending request for text = Studies of the furious and paralytic forms of canine rabi es at the early stage of disease have shown Pausing for 10 secs... OK Sending request for text = Replication fork integrity , which is essential for the m aintenance of genome stability , is monitor Pausing for 10 secs... OK Sending request for text = Polyarthritis and rash caused by Sindbis virus ( SINV ) , was first recognised in northern Europe ab Pausing for 10 secs... OK Sending request for text = With the post-genomic era came a dramatic increase in hig h-throughput technologies , of which transc Pausing for 10 secs... OK Sending request for text = Centromeres are the attachment points between the genome and the cytoskeleton: centromeres bind to k Pausing for 10 secs... OK Sending request for text = Humans are a diploid species that inherit one set of chro mosomes paternally and one homologous set o Pausing for 10 secs... Sending request for text = In contrast to HIV infection in humans and SIV in macaque s , SIV infection of natural hosts includin Pausing for 10 secs... Sending request for text = Individuals choose their mates so as to maximize reproduc tive success , and one important component Pausing for 10 secs... OK Sending request for text = Breast cancers that are "triple-negative" for the clinica 1 markers ESR1 , PGR , and HER2 typically b Pausing for 10 secs... Sending request for text = Pathogen-associated secretion systems translocate numerou s effector proteins into eukaryotic host ce Pausing for 10 secs... OK Sending request for text = The inference of regulatory interactions and quantitative models of gene regulation from time-series Pausing for 10 secs... Sending request for text = Soil-transmitted helminth ( STH ) infections ( i . e . , Ascaris lumbricoides , hookworm , and Trich Pausing for 10 secs... Sending request for text = Prompt post-exposure prophylaxis ( PEP ) is essential in preventing the fatal onset of disease in pe Pausing for 10 secs... OK Sending request for text = Uganda has active foci of both chronic and acute HAT with the acute zoonotic form of disease classic

OK Sending request for text = Dietary restriction ( DR ) extends lifespan in various sp ecies and also slows the onset of age-relat Pausing for 10 secs... OK Sending request for text = The unique capability of acetogens to ferment a broad ran ge of substrates renders them ideal candida Pausing for 10 secs... Sending request for text = Arabidopsis thaliana cryptochrome 2 ( CRY2 ) mediates lig ht control of flowering time . CIB1 ( CRY2-Pausing for 10 secs... OK Sending request for text = Enteric bacterial pathogens cause food borne disease , wh ich constitutes an enormous economic and he Pausing for 10 secs... OK Sending request for text = High-altitude hypoxia ( reduced inspired oxygen tension d ue to decreased barometric pressure ) exert Pausing for 10 secs... OK Sending request for text = Machupo virus ( MACV ) , a New World arenavirus , is the etiological agent of Bolivian hemorrhagic f Pausing for 10 secs... Sending request for text = Serological tests for IgM and IgG are routinely used in c linical laboratories for the rapid diagnosi Pausing for 10 secs... Sending request for text = Toll/interleukin-1 receptor ( TIR ) domains in Toll-like receptors are essential for initiating and Pausing for 10 secs... OK Sending request for text = Protein modifications play a major role for most biologic al processes in living organisms . Amino-te Pausing for 10 secs... Sending request for text = Exosomes can transfer genetic materials between cells . T heir roles in viral infections are beginnin Pausing for 10 secs... OK Sending request for text = According to recent experimental evidence , the interacti on between chromatin loops , which can be c Pausing for 10 secs... Sending request for text = Predicting the dynamic behavior of a large network from t hat of the composing modules is a central p Pausing for 10 secs... Sending request for text = The 5-year survival of non-small cell lung cancer patient s can be as low as 1% in advanced stages . Pausing for 10 secs... OK Sending request for text = The mechanisms and treatment of psychomotor retardation ,

which includes motor and cognitive impairm

OK Sending request for text = Natural selection drives populations towards higher fitne ss, but crossing fitness valleys or platea Pausing for 10 secs... OK Sending request for text = The nuclear pore complex ( NPC ) regulates molecular traf fic across the nuclear envelope ( NE ) . Se Pausing for 10 secs... OK Sending request for text = Staphylococcus aureus is an opportunistic pathogen that c olonizes the skin and mucosal surfaces of m Pausing for 10 secs... OK Sending request for text = The Type II Secretion System ( T2SS ) is a molecular mach ine that drives the secretion of fully-fold Pausing for 10 secs... OK Sending request for text = Burkholderia pseudomallei is a mostly saprophytic bacteri um , but can infect humans where it causes Pausing for 10 secs... OK Sending request for text = Offspring of Schistosoma mansoni-infected women in schist osomiasis-endemic areas may be sensitised i Pausing for 10 secs... Sending request for text = Hox proteins play fundamental roles in controlling morpho genetic diversity along the anterior-poster Pausing for 10 secs... Sending request for text = The antiproliferative response to anticancer treatment is the result of concurrent responses in all Pausing for 10 secs... OK Sending request for text = Horizontal acquisition of DNA by bacteria dramatically in creases genetic diversity and hence success Pausing for 10 secs... Sending request for text = HIV-1 vaccines designed to date have failed to elicit neu tralizing antibodies ( Nabs ) that are capa Pausing for 10 secs... OK Sending request for text = Approaches based on linear mixed models ( LMMs ) have rec ently gained popularity for modelling popul Pausing for 10 secs... Sending request for text = Transient associations among neurons are thought to under lie memory and behavior . However , little Pausing for 10 secs... OK Sending request for text = Polarized growth is maintained by both polarized exocytos is , which transports membrane components t Pausing for 10 secs...

OK

Sending request for text = Miltefosine , an anti-cancer drug that has been successfully repositioned for treatment of Leishmani Pausing for 10 secs...

OK Sending request for text = In neurons polarized trafficking of vesicle-bound membran e proteins gives rise to the distinct molec Pausing for 10 secs... OK Sending request for text = The consensus that complexity begets stability in ecosyst ems was challenged in the seventies , a res Pausing for 10 secs... Sending request for text = Vibrio cholerae is a bacterial pathogen that colonizes th e chitinous exoskeleton of zooplankton as w Pausing for 10 secs... OK Sending request for text = Marburg virus , the Kaposi's sarcoma-associated herpesvir us ( KSHV ) and Dengue virus all activate, Pausing for 10 secs... OK Sending request for text = The development of new drugs against Chagas disease is a priority since the currently available medi Pausing for 10 secs... OK Sending request for text = Bistability plays a central role in the gene regulatory n etworks ( GRNs ) controlling many essential Pausing for 10 secs... Sending request for text = Even in the absence of an adaptive immune system in murin e models , lymphatic dilatation and dysfunc Pausing for 10 secs... Sending request for text = Plant-pathogenic Xanthomonas bacteria secrete transcripti on activator-like effectors ( TALEs ) into Pausing for 10 secs... OK Sending request for text = The evolution of new gene networks is a primary source of genetic innovation that allows bacteria to Pausing for 10 secs... Sending request for text = Two-component signaling systems are ubiquitous in bacteri a , Archaea and plants and play important r Pausing for 10 secs... OK Sending request for text = The bacterial replication cycle is driven by the DnaA pro tein which cycles between the active ATP-bo Pausing for 10 secs... Sending request for text = The rat demyelination ( dmy ) mutation serves as a unique model system to investigate the maintenanc Pausing for 10 secs... Sending request for text = Leishmaniasis is an important disease that affects 12 mil lion people in 88 countries , with 2 millio Pausing for 10 secs...

OK

Sending request for text = To gain a more detailed picture of cryptococcosis in Thai land , a retrospective study of 498 C . neo Pausing for 10 secs...

OK Sending request for text = Scabies is one of the commonest dermatological conditions globally; however it is a largely underexp Pausing for 10 secs... OK Sending request for text = The type III interferon ( IFNλ ) receptor IL-28R is abund antly expressed in the respiratory tract an Pausing for 10 secs... OK Sending request for text = Neural networks with a single plastic layer employing rew ard modulated spike time dependent plastici Pausing for 10 secs... OK Sending request for text = Genome-wide association studies ( GWAS ) have been fruitf ul in identifying disease susceptibility lo Pausing for 10 secs... OK Sending request for text = Leishmaniasis is endemic in 98 countries with an estimate d 350 million people at risk and approximat Pausing for 10 secs... OK Sending request for text = IKAROS is a critical regulator of hematopoietic cell fate and its dynamic expression pattern is requ Pausing for 10 secs... Sending request for text = Mitochondria originated from proteobacterial endosymbiont s , and their transition to organelles was Pausing for 10 secs... Sending request for text = Genetic factors play an important role in the etiology of both sporadic and familial breast cancer . Pausing for 10 secs... OK Sending request for text = Understanding complex networks of protein-protein interac tions ( PPIs ) is one of the foremost chall Pausing for 10 secs... Sending request for text = Buruli ulcer ( BU ) , caused by infection with Mycobacter ium ulcerans , is a chronic necrotizing hum Pausing for 10 secs... OK Sending request for text = The mosquito Aedes aegypti , vector of dengue , chikungun ya and yellow fever viruses , is an importa Pausing for 10 secs... Sending request for text = Mean-field approximations are a powerful tool for studyin g large neural networks . However , they do Pausing for 10 secs... Sending request for text = A major problem in biology is to understand how complex t issue shapes may arise through growth . In Pausing for 10 secs... OK Sending request for text = Ophthalmo-acromelic syndrome ( OAS ) , also known as Waar denburg Anophthalmia syndrome , is defined

OK Sending request for text = The Drosophila genes spalt major ( salm ) and spalt-relat ed ( salr ) encode Zn-finger transcription Pausing for 10 secs... OK Sending request for text = Sensory systems adapt their neural code to changes in the sensory environment, often on multiple ti Pausing for 10 secs... OK Sending request for text = While the roles of rpoSBb and RpoS-dependent genes have b een studied extensively within the mammal, Pausing for 10 secs... OK Sending request for text = Two-cysteine peroxiredoxins are ubiquitous peroxidases th at play various functions in cells . In Lei Pausing for 10 secs... OK Sending request for text = Microbial metabolism of plant polysaccharides is an impor tant part of environmental carbon cycling, Pausing for 10 secs... OK Sending request for text = Glanders , caused by the gram-negative bacterium Burkhold eria mallei , is a highly infectious zoonot Pausing for 10 secs... Sending request for text = Observations gained from model organisms are essential, yet it remains unclear to which degree they Pausing for 10 secs... Sending request for text = The 3D organization of chromosomes is crucial for regulat ing gene expression and cell function . Man Pausing for 10 secs... OK Sending request for text = Toxoplasma is an obligate intracellular parasite that rep licates in mammalian cells within a parasit Pausing for 10 secs... Sending request for text = The BMP signaling pathway has a conserved role in dorsalventral axis patterning during embryonic de Pausing for 10 secs... OK Sending request for text = The mitochondrial protein repertoire varies depending on the cellular state . Protein component modi Pausing for 10 secs... Sending request for text = The dissociation mechanism of the thioredoxin ( Trx ) mix ed disulfide complexes is unknown and has b Pausing for 10 secs... Sending request for text = Pathophysiological mechanisms are still incompletely unde rstood for leprosy , an urgent public healt Pausing for 10 secs... OK Sending request for text = Metagenomic sequencing of patient samples is a very promi sing method for the diagnosis of human infe

OK Sending request for text = Environmental exposures filtered through the genetic make -up of each individual alter the transcript Pausing for 10 secs... OK Sending request for text = Every instant of perception depends on a cascade of brain processes calibrated to the history of sen Pausing for 10 secs... OK Sending request for text = Examples of animals evolving similar traits despite the a bsence of that trait in the last common and Pausing for 10 secs... OK Sending request for text = Human papillomaviruses ( HPVs ) are DNA viruses associate d with major human cancers . As such there Pausing for 10 secs... OK Sending request for text = Orofacial clefting is amongst the most common of birth de fects , with both genetic and environmental Pausing for 10 secs... OK Sending request for text = The mechanisms of hypoxic injury to the developing human brain are poorly understood , despite being Pausing for 10 secs... Sending request for text = Filarial nematodes maintain a mutualistic relationship wi th the endosymbiont Wolbachia . Depletion o Pausing for 10 secs... Sending request for text = Genetic reassortment between influenza A viruses ( IAVs ) facilitate emergence of pandemic strains, Pausing for 10 secs... OK Sending request for text = The Saccharomyces cerevisiae polo-like kinase Cdc5 promot es adaptation to the DNA damage checkpoint Pausing for 10 secs... Sending request for text = Sequences of higher frequency A and lower frequency B ton es repeating in an ABA- triplet pattern are Pausing for 10 secs... OK Sending request for text = Recent interest in male-based sterile insect technique ( SIT ) and incompatible insect technique ( I Pausing for 10 secs... Sending request for text = Hematophagous mosquitos and ticks avoid host hemostatic s ystem through expression of enzyme inhibito Pausing for 10 secs... Sending request for text = In mammals , dosage compensation is achieved by doubling expression of X-linked genes in both sexes Pausing for 10 secs... OK Sending request for text = The opportunistic human fungal pathogen , Candida albican

s , undergoes morphological and transcripti

OK Sending request for text = Egress of newly assembled herpesvirus particles from infe cted cells is a highly dynamic process invo Pausing for 10 secs... OK Sending request for text = The variants of human influenza virus have caused , and c ontinue to cause , substantial morbidity an Pausing for 10 secs... Sending request for text = Protozoan parasites of the genus Leishmania alternate bet ween flagellated , elongated extracellular Pausing for 10 secs... OK Sending request for text = To further investigate the importance of insulin signalin g in the growth , development , sexual matu Pausing for 10 secs... OK Sending request for text = Primate lentivirus nef is required for sustained virus re plication in vivo and accelerated progressi Pausing for 10 secs... OK Sending request for text = The RNA polymerase II ( Pol II ) is a eukaryotic enzyme t hat catalyzes the synthesis of the messenge Pausing for 10 secs... Sending request for text = Visual object recognition and sensitivity to image featur es are largely influenced by contextual inp Pausing for 10 secs... Sending request for text = Lectin-like bacteriotoxic proteins , identified in severa l plant-associated bacteria, are able to s Pausing for 10 secs... OK Sending request for text = The genomic GC-content of bacteria varies dramatically, from less than 20% to more than 70% . This Pausing for 10 secs... Sending request for text = The expansion of CAG/CTG repeats is responsible for many diseases , including Huntington's disease ( Pausing for 10 secs... OK Sending request for text = Research on Neglected Tropical Diseases ( NTDs ) has incr eased in recent decades , and significant n Pausing for 10 secs... Sending request for text = Chagas disease is the most important vector-borne disease in Latin America . Regional initiatives ba Pausing for 10 secs... Sending request for text = The weight with which a specific outcome feature contribu tes to preference quantifies a person's 'ta Pausing for 10 secs... OK Sending request for text = Rice is an important monocotyledonous crop worldwide; it differs from the dicotyledonous plant Arabi

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OK
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ssues and its elevated expression is associ Pausing for 10 secs... OK Sending request for text = The majority of HIV-1 infections in women occur through v aginal intercourse , in which virus-contain Pausing for 10 secs... Sending request for text = The evolutionary origins of genetic robustness are still under debate: it may arise as a consequence Pausing for 10 secs... OK Sending request for text = Histone chaperones CAF-1 and Asf1 function to deposit new ly synthesized histones onto replicating DN Pausing for 10 secs... OK Sending request for text = Vaccination is an effective method to protect against inf ectious diseases . An important considerati Pausing for 10 secs... OK Sending request for text = Changes in genomic DNA methylation patterns are generally assumed to play an important role in the e Pausing for 10 secs... Sending request for text = The PR interval on the electrocardiogram reflects atrial and atrioventricular nodal conduction time Pausing for 10 secs... Sending request for text = Inhibitory interneurons play critical roles in shaping th e firing patterns of principal neurons in m Pausing for 10 secs... OK Sending request for text = The BRCA Challenge is a long-term data-sharing project in itiated within the Global Alliance for Geno Pausing for 10 secs... Sending request for text = The small GTPase RAS is among the most prevalent oncogene s . The evolutionarily conserved RAF-MEK-MA Pausing for 10 secs... OK Sending request for text = Recent years have seen the development of numerous method ologies for reconstructing transmission tre Pausing for 10 secs... Sending request for text = Hyperendemic circulation of all four types of dengue viru s ( DENV-1-4 ) has expanded globally , fuel Pausing for 10 secs... OK Completed In [28]: # check how many rows have blank result (some errors) empty\_df = df.query("groq\_mistral\_summary.str.strip() == ''") print(len(empty df))

Sending request for text = The long noncoding MALAT1 RNA is upregulated in cancer ti

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In [29]: # attempt to retry for blank rows (due to some errors)
         retry = True
         if retry:
             print("Retrying for empty results...")
             for i in range(len(df)):
                 item = df.iloc[i]
                 if item["groq_mistral_summary"] == "":
                     print("Item =", i)
                     text = item["article"]
                     df.at[i, "groq_mistral_summary"] = send_sumarize_request(text[:text_cap
             print("Completed")
```

Retrying for empty results... Completed

df.head()							
	lay_summary	article	headings	keywords	id	gr	
0	Yersinia pestis , the bacterial agent of plagu	Fleas can transmit Yersinia pestis by two mech	[Abstract, Introduction, Results, Discussion, 	[united states, invertebrates, medicine and he	journal.ppat.1006859		
1	The genome of all vertebrates is heavily colon	Endogenous retroviruses ( ERVs ) are remnants	[Abstract, Introduction, Results, Discussion, 	[viruses, sheep, virology]	journal.ppat.0030170		
2	The molecular mechanisms underlying directed c	The Drosophila embryonic gonad is assembled fr	[Abstract, Introduction, Results, Discussion, 	0	journal.pgen.1003720		
\$	Contrary to the long- standing belief that no n	Recently , we presented a study of adult neuro	[Abstract, Introduction, Model, Results, Discu	[computational biology/computational neuroscie	journal.pcbi.1001063	re	
ı	Embryonic stem cells have two remarkable prope	Understanding the transcriptional regulation o	[Abstract, Introduction, Results, Discussion, 	[developmental biology, cell biology, mammals,	journal.pgen.0030145		
	0	lay_summary  Yersinia pestis , the bacterial agent of plagu  The genome of all vertebrates is heavily colon  The molecular mechanisms underlying directed c  Contrary to the long-standing belief that no n  Embryonic stem cells have two remarkable	Yersinia pestis , the bacterial agent of plagu  The genome of all vertebrates is heavily colon  The molecular mechanisms underlying directed c  Contrary to the long-standing belief that no n  Embryonic stem cells have two remarkable regulation o	Yersinia pestis, the bacterial agent of plagu  The genome of all vertebrates is heavily colon  The molecular mechanisms underlying directed c  Contrary to the longstanding belief that no n  Embryonic stem cells have two remarkable regulation o  Yersinia pestis transmit Yersinia pestis by two mech  Endogenous retroviruses (ERVs ) are retroviruses (ERVs ) are remnants  [Abstract, Introduction, Results, Discussion, Introduction, Results, Discussion, Introduction, Results, Discussion, Introduction, Results, Discussion, Introduction, Introduction, Introduction, Model, Results, Discussion, Introduction, Results, Discussion, Dis	Persinia pestis   Fleas can transmit   Agent of plagu   The genome of all vertebrates is heavily colon   ERVs ) are remnants   Contrary to the long-belief that no n   Computational stamp belief that no remarkable   Embryonic stem cells have two remarkable   Line plagus   Line plagus   Line plagus   Fleas can transmit tr	Versinia pestis   Fleas can transmit yersinia pestis   Fleas can transmit yersinia pestis   Plagu   Introduction, Results, Discussion,   Introduction, Results, Discussion,	

```
output_filename = "plos_groq_mistral_summary.csv"
print("Writing to file ", output_filename)
df.to_csv(output_path+output_filename,
          index = False
```