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# The Arcuate Fasciculus and language origins: *Disentangling existing conceptions that influence evolutionary accounts*

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## Abstract

*The Arcuate Fasciculus (AF) is of considerable interdisciplinary interest, because of its major implication in language processing. Theories about language brain evolution are based on anatomical differences in the AF across primates. However, changing methodologies and nomenclatures have resulted in conflicting findings regarding interspecies AF differences: Historical knowledge about the AF originated from human blunt dissections and later from monkey tract-tracing studies. Contemporary tractography studies reinvestigate the fasciculus' morphology, but remain heavily bound to unclear anatomical priors and methodological limitations. First, we aim to disentangle the influences of these three epistemological steps on existing AF conceptions, and to propose a contemporary model to guide future work. Second, considering the influence of various AF conceptions, we discuss four key evolutionary changes that propagated current views about language evolution: 1) frontal terminations, 2) temporal terminations, 3) greater Dorsal- versus Ventral Pathway expansion, 4) lateralisation. We conclude that new data point towards a more shared AF anatomy across primates than previously described. Language evolution theories should incorporate this continuous AF evolution across primates.*

**Keywords:** language network, language evolution, frontal terminations, temporal terminations, discontinuity, continuity, lateralization, nomenclature, terminology, hemispheric specialization, comparative anatomy, white matter, tractography, evolutionary neuroscience, monkey, nonhuman primate

**Abbreviations:** AF (Arcuate Fasciculus), SLF (Superior Longitudinal Fasciculus), MdLF (Middle Longitudinal Fasciculus), dMRI (diffusion magnetic resonance imaging), NHP (Nonhuman primate)

## Highlights:

- 1) Existing conceptions of the AF are disentangled to derive a current model
- 2) Data indicates a continuity in AF's frontal terminations across primates
- 3) Data indicates a possible continuity in AF's temporal terminations across primates
- 4) Data indicates a possible continuity in the Dorsal- versus Ventral Pathways strength
- 5) More data is needed to conclude about the (dis)continuity of AF's lateralisation

## Introduction

The Arcuate Fasciculus (AF), or the *arched/curved bundle* in Latin, is traditionally regarded as one of the major language pathways in the human brain: It arches around the Sylvian or lateral fissure to connect the key speech production region (Broca's area) in the frontal lobe, with the speech comprehension region (Wernicke's area) in the posterior temporal lobe (Geschwind 1970). Congruent with Geschwind's influential model, decades of subsequent research have generally converged on the AF's critical involvement in language and speech functions (e.g., Catani and Mesulam, 2008; Catani and Schotten, 2012; Dick and Tremblay, 2012). Interestingly, the anatomical homologue of this critical white matter pathway for human language exists in the nonhuman primates, where language is absent (e.g., Eichert et al. 2019; Frey et al. 2014; Rilling et al. 2008, 2012). This crucial phylogenetic link provides a valuable avenue to probe the evolutionary origins of human language i.e., by comparing how the AF differs across human and nonhuman primates. Indeed, in the past decade, the emergent field of nonhuman primate or comparative magnetic resonance imaging (MRI) neuroimaging (Friedrich et al. 2021; Milham et al. 2020) have enabled detailed anatomical and functional comparisons of the AF across humans and various nonhuman primate species, such as the macaque monkey and the chimpanzee (e.g., Balezeau et al. 2020; Mars et al., 2021; Rilling et al. 2008, 2012; Rocchi et al. 2021; Wilson et al. 2015). The overarching aim of this review is to synthesize this exciting new body of work that compares the anatomy and function of the AF in humans versus nonhuman primates, to understand *how the Arcuate Fasciculus might, or might not, have changed across primate evolution to support the emergence of language in humans.*

In the first part of this review, we tackle a major challenge that confronts present-day researchers seeking to investigate the AF: *There is currently poor consensus about the anatomical definition of the AF* (e.g., where it terminates in the frontal and temporal lobes), *and its nomenclature* (e.g., whether it is considered part of, or distinct from the Superior Longitudinal Fasciculus (SLF) or Middle Longitudinal Fasciculus (MdLF)). This is primarily driven by the fact that, since its discovery, the methodologies used to study and delineate the AF (e.g., blunt dissections, diffusion MRI tractography, neural tract-tracing) have been constantly evolving, resulting in different descriptions and naming of the AF (e.g., Oliveira et al., 2021; Vavassori et al., 2021). As such, we start the current review with an overview of how the definition and nomenclature of the AF have evolved along with the emergence of different methodologies in the field. This overview, in disentangling the various existing conceptions of the AF, aims to provide a contemporary definition of the fibre bundle that will guide our subsequent review of existing comparative studies of the primate AF.

The second part of this review will discuss four key findings from recent comparative research that shed light on the question about how the AF has changed in humans relative to nonhuman primates: 1) *An expansion of the AF's frontal terminations into the inferior frontal cortex* (e.g., Schmahmann and Pandya, 2006); 2) *An expansion of the AF's temporal terminations beyond the middle temporal cortex* (e.g., Eichert et al. 2019); 3) *A greater expansion of the Dorsal Language Pathway (which includes the AF), relative to the Ventral Language Pathway* (e.g., Rilling et al., 2008); and 4) *An absence of lateralisation in the monkey's AF* (e.g., Balezeau et al., 2020). For each finding, we will discuss how the different conceptions of the AF have led to different views on these questions, and subsequently influenced theorisations about the evolution of language. We will conclude that the collective data point towards a continuity in AF anatomy across primates. Last but not least, this review will open a discussion regarding the existing questions and future directions in the field.

## Part I: Disentangling existing conceptions of the Arcuate Fasciculus

Differing anatomical descriptions and nomenclatures of the Arcuate Fasciculus (AF) fibre bundle have surfaced since its initial discovery by Reil in 1809 (see Catani et al., 2010; Dick and Tremblay 2012; ef. Oliveira et al., 2021; Vavassori et al., 2021 for historical reviews of the AF). To a large extent, the various conceptions result from the various emerging methodologies used to investigate the AF, including post-mortem brain dissections, neural tract-tracing, and more recently, diffusion magnetic resonance imaging (dMRI) tractography. Here, we briefly trace the emergence of various conceptions of the AF, relating them to the methods employed to investigate the tract, to finally arrive at a contemporary definition of the AF to guide our subsequent review of the comparative AF literature.

### Early 19<sup>th</sup> to late 20<sup>th</sup> century: Post-mortem blunt dissections and the synonymity between Arcuate Fasciculus (AF) and Superior Longitudinal Fasciculus (SLF)

*Post-mortem blunt dissection* was the first methodology adopted to investigate the Arcuate Fasciculus (AF). The method briefly involves first, the post-mortem extraction and fixation of the brain, followed by the use of a blunt dissection tool to meticulously peel away the grey and white matter progressively to reveal the extents of the various axon bundles. The blunt dissection method was the dominant method employed throughout the 19<sup>th</sup> century to study human neuroanatomy and had led to many seminal works and discoveries of that era (e.g., Burdach 1826; Dejerine, Dejerine, and Dejerine 1895; 1901; Gratiolet, 1839; Meynert, 1885; Klingler, 1935). While this technique enabled the identification of distinct white matter bundles and their individual courses in the brain, they were limited in 1) making distinctions between axon bundles that run in parallel, or that overlap or cross in their courses; and 2) determining the precise axonal origins and terminations of the bundles (Petrides 2014; Yaşargil et al. 2004).

Indeed, it was through blunt dissections that Johann Christen Reil, in 1809, first discovered the system of longitudinal white matter fibres that arched along the Sylvian (or lateral) fissure to interconnect the frontal, parietal and temporal cortices (Figure 1). Using the same method, Friedrich Burdach further characterised and first named these long arching fibres as the “Bogenbündel” in German, which translates to the “arched bundle” or *arcuate fasciculus* in Latin (Burdach, 1822). For most of the 19<sup>th</sup> century, two nomenclatures -- “Arcuate Fasciculus (AF)” and “Superior Longitudinal Fasciculus (SLF)” were synonymously used by the most prominent neuroanatomists to describe the longitudinal frontal-parieto-temporal fibre bundle (Catani et al., 2010; Dick and Tremblay, 2012). For instance, while Meynert (1885) used the term AF, Wernicke (1874) and Déjérine (1895, 1901) used both AF and SLF to refer to the same longitudinal fibre bundle. The interest for the SLF/AF fibre bundle rose to prominence at the start of the 20<sup>th</sup> century as Wernicke (1908) identified it as one of two main associative fibre systems for language. Notably, in Geschwind's influential neurological model of language (1970), the term AF, and not the term SLF, was used to depict the fibre system linking the language production centre (inferior frontal cortex) to the language comprehension centre (posterior middle temporal cortex). Since then, the label AF had received an increased recognition and usage to refer to the classical language pathway (more than the term SLF), even though the terms AF and SLF are generally referred to the same tract bundle until the late 20<sup>th</sup> century (Figure 1).

Figure 1: Classic dissection derived AF/SLF Model (Reil, 1809; Burdach, 1822; Déjèrine, 1895)

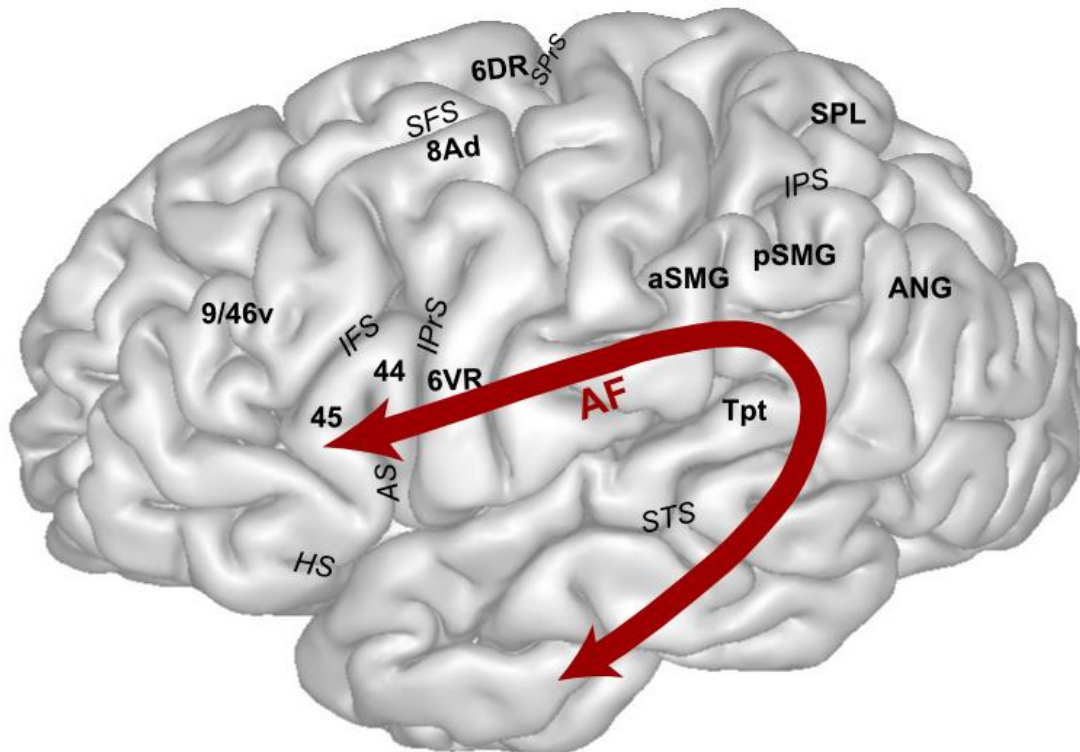


Figure 1: Classic dissection derived AF/SLF Model (e.g., Reil, 1809; Burdach, 1822; Déjèrine, 1895, 1901) in the human brain. AF: Arcuate Fasciculus; SFS=Superior Frontal Sulcus; SPrS=Superior Precentral Sulcus; IPrS=Inferior Precentral Sulcus; IFS=Inferior Frontal Sulcus; AS=Ascending branch of Sylvian Fissure; HS=Horizontal branch of Sylvian Fissure; IPS=Intra-Parietal Sulcus; Tpt=Temporal-parietal area (BA22); aSMG=anterior Supra-Marginal Gyrus (BA40); pSMG=posterior Supra-Marginal Gyrus (BA40); ANG=Angular Gyrus (BA39); SPL=Superior Parietal Lobule (BA7).

From the turn of the 20th century until today, two other methodologies were adopted, alongside blunt dissections (see Dziedzic et al., 2021 for a recent review on dissections with the Klingler technique), to investigate the Arcuate Fasciculus bundle: *neural tract-tracing* and *diffusion MRI (dMRI) tractography*. These techniques have led to dissociate the classic AF/SLF longitudinal fibre bundle into different constituent bundles with distinct courses and terminations. However, due to the differences between the two methods, various models of the AF/SLF fibre system had emerged. Here, we briefly describe each method and its associated models and conceptions of the AF/SLF fibre system.

### Late 20th Century: Neural tract-tracing reveals distinctions between the AF and SLF fibre systems

1        *Neural tract-tracing* is a precise, but also highly invasive method for studying brain  
2 connections (see Lanciego and Wouterlood, 2020; Saleeba et al., 2019 for recent reviews about  
3 the tract-tracing methodology). Briefly, it involves the injection of anatomical tracers into a living  
4 brain, which would be transported from the injected region to other regions via interconnecting  
5 axons, followed by histological analyses of the brain ex-vivo to visualise the axons and their  
6 terminations that have been labelled by the injected tracers. Depending on whether a *retrograde*  
7 or *anterograde* tracer was administered, the brain region that *sends projections into* the injected  
8 region or the brain region that *receives projections from* the injected region along with the  
9 interconnecting axons, will be labelled, respectively. Hence, using a combination of retrograde and  
10 anterograde tracing, the origins and terminations of cortico-cortical connections associated with a  
11 brain region of interest can be precisely determined. As such, contrary to blunt dissection and  
12 dMRI tractography, tract-tracing possesses the resolution to distinguish between fibre bundles that  
13 overlap or cross in their courses but have different terminations and can therefore also define a  
14 monosynaptical connection. Thus, it is commonly regarded as the gold-standard technique for  
15 studying brain connections. A key drawback, however, is that tract-tracing is currently only  
16 performed in few nonhuman primate subjects due to its invasive nature, which requires the  
17 euthanasia of the animal.

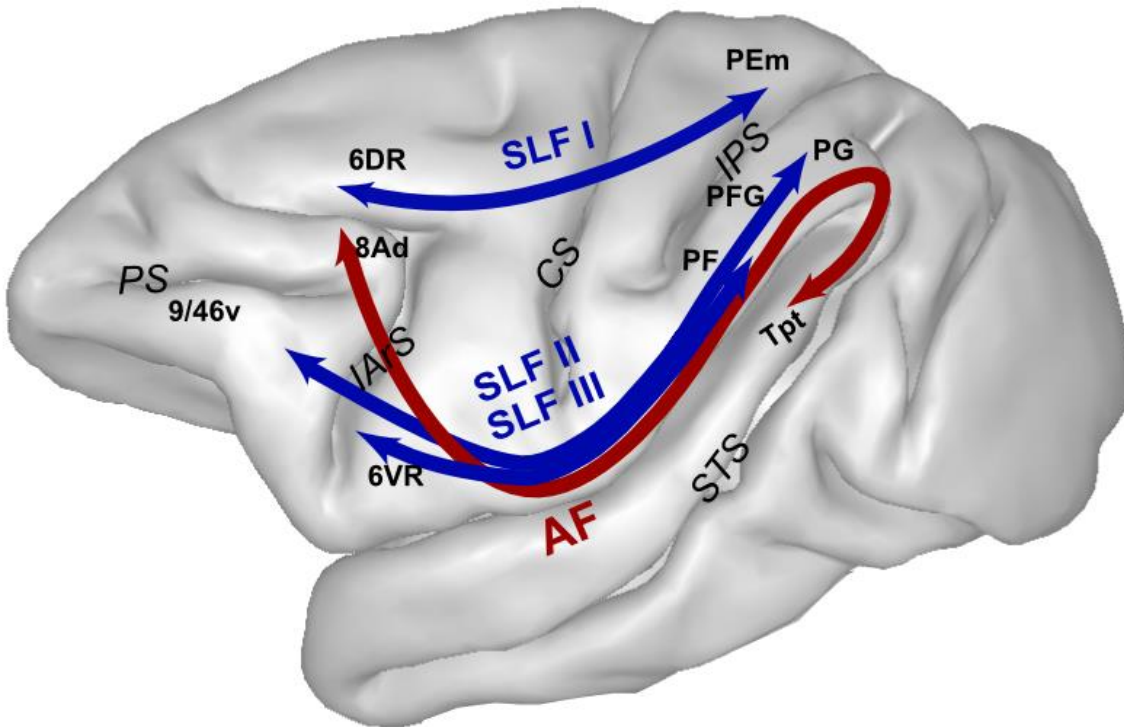
18        Tract-tracing was employed in a series of important studies that resulted in the dissociation  
19 of the classical AF/SLF longitudinal fibre system into several distinct fibre bundles based on their  
20 origins and terminations in the macaque brain (Frey et al., 2014; Petrides and Pandya, 1984, 1988,  
21 2006, 2009; Schmahmann et al., 2007; Schmahmann and Pandya, 2006; Yeterian et al., 2012 etc.)  
22 (Figure 2). In these studies, anatomical tracers were injected into various frontal, parietal and  
23 temporal regions in the macaque brain, which have known homologies with the human brain, to  
24 study their interconnections. Notably, this body of work has led to a new model of the AF/SLF  
25 longitudinal fibre system which consists of four distinct fibre bundles including:

26        1) the AF, which comprise the fibres that arch dorsally around the Sylvian fissure and  
27 connect the posterior temporal cortex with the frontal cortex (the exact frontal and temporal  
28 terminations of the AF will be further discussed in Part 2, see also Figure 5); and

29        2) the SLF I, II and III, which comprise the three longitudinal fibre bundles that  
30 interconnect distinct parts of the frontal and parietal cortex and are not arching around the Sylvian  
31 fissure (Figure 2).

32        Since the advent of this approach, the term SLF is used to designate portions of the AF/SLF  
33 complex, which are distinct from the AF and which are further distinguished in three different  
34 branches I, II and III (Figure 2).

1 Figure 2: Initial Macaque AF and SLF Tract-tracing Model (Petrides and Pandya, 1988,  
2 Schmahmann and Pandya, 2006)



3 Figure 2: Initial Tract-tracing Model that separated the AF from the SLF in the macaque brain.  
4 Adapted from Petrides and Pandya (1988), Schmahmann and Pandya (2006). SLF I, II, III=First,  
5 second and third branch of the Superior Longitudinal Fasciculus, AF=Arcuate Fasciculus;  
6 IArS=Inferior Arcuate Sulcus, PS=Principal Sulcus; CS=Central Sulcus; IPS=Intra-Parietal  
7 Suclus; ; STS=Superior Temporal Sulcus; PF; PEm; PFG; PG are cytoarchitectonic labels for  
8 the parietal areas based on Von Economo and Koskinas (1925)

### 11 Towards a contemporary conception of the AF: diffusion MRI tractography

13 *Diffusion MRI (dMRI) tractography* is, at present, the only technique available to  
14 investigate white matter fibre pathways both in-vivo and ex-vivo, across a large sample of  
15 individuals. It involves a form of magnetic resonance imaging that measures the diffusivity of  
16 water molecules in the brain. Since water diffusion in the brain is highly constrained by axons (i.e.,  
17 water tends to flow along the direction of the axons), by tracking the most probable directions of  
18 water diffusion voxel-by-voxel, one would be able to reconstruct the main fibre pathways that  
19 course between two specific areas in the brain. Alternatively, even the total pathways between all  
20 areas in the brain can be reconstructed, yielding a so-called *whole-brain tractogram*. Note that  
21 these reconstructed pathways are essentially “streamlines” that connect vectors of water diffusion  
22 computed in each brain voxel and might not necessarily depict true axonal connections. Therefore,  
23 tractograms notoriously contain a large number of false positive (Maier-Hein et al. 2017). This is  
24 especially so when there are multiple crossing fibres passing within a single voxel, which makes

it notoriously difficult to determine the actual orientation of fibres (Girard et al. 2020). Additionally, dMRI tractography does not reveal whether a connection is monosynaptic (single axon from origin to termination sites) or polysynaptic (more than one axon interconnecting between origin and termination sites). Due to the limitations in disentangling the various association pathways that pass through the frontal and parietal-temporal regions (i.e., AF, SLF I, II, III), most dMRI studies therefore collectively refer to these tracts as the “Dorsal language Pathway or stream” (e.g., Saur et al., 2008). On the other hand, a second set of pathways that curves ventrally from the inferior prefrontal cortex, across the Sylvian fissure, to the anterior and middle temporal cortex are collectively known as the “Ventral language Pathway or stream”. This division emulates the well-known visual pathway division (Hickok and Poeppel, 2004). Functionally, the Dorsal Pathway is involved in mapping sound to articulation, while the Ventral Pathway is involved in mapping sound to meaning (Hickok and Poeppel, 2004; Saur et al., 2008). However recent advances are challenging this dichotomic view and are also highlighting the importance of the Dorsal Pathway in semantics (Hula et al. 2020). Note that the current review is centred around the Dorsal Pathways and the AF’s definition and its implication in brain and language evolution. Going into the debate of a Ventral Pathway’s definition would exceed the current review. We will report exclusively data concerning the proportional strength between the Dorsal and Ventral Pathways, that are further discussed in the following section (Question 4, Figure 5).

Here we discuss two contemporary and influential conceptions of the classical SLF/AF bundle and how they were derived based on different theoretical underpinnings:

The first conception, which was proposed by Catani and colleagues using dMRI tractography (Catani et al., 2005), is based on priors from classical dissection studies (Figure 3 left panel). This conception names the peri-sylvian longitudinal fibres (both arching and non-arching) collectively as the AF, and divides it into three distinct segments: 1) a *long direct segment* that connects the posterior part of the superior temporal gyrus (Wernicke’s area) to the inferior frontal cortex (Broca’s area); 2) an *anterior indirect segment* that connects Broca’s area and inferior (Geschwind’s territory) parietal cortices; and 3) a posterior indirect segment that links the inferior parietal cortices with the Wernicke’s area. This model was obtained via a two-regions-of-interest (ROI) “virtual dissection” approach using diffusion tractography (Catani et al., 2002). First, one ROI was defined that delineated the main body of the AF identified on the fractional anisotropy (FA) image. Next, two other ROIs were defined in the inferior frontal cortex and posterior superior temporal cortex respectively. Fibres were then tracked between each pair of ROIs which yielded the three segments of the AF. Note that the placement of ROIs was strongly influenced by Geschwind’s model of the language system (1970) that proposed a pathway, which was termed the AF, linking Broca’s and Wernicke’s language areas in the brain. Indeed, this 3-segment AF model was consistent with neurological models of aphasia and evidence from aphasic patients (e.g., Damasio and Geschwind 1984, Lichtheim, 1885), where the *direct long pathway* (frontal-temporal) is associated with *phonological-related language processes* (e.g., rapid word repetitions that do not involve semantic retrieval processes); while the *indirect anterior* (frontal-parietal) and *posterior* (parietal-temporal) pathways are associated with *semantic-related language processes*, such as speech comprehension and the production of meaningful words or speech, respectively (Catani et al., 2005). Note that the tractography recipe of that model has been improved in recent years, including more than two ROIs (e.g., Forkel et al., 2014). Subsequently, this 3-segment AF model was also validated by other studies using human electrocorticography (Matsumoto et al.



2004), functional connectivity (Holland et al., 2007) and post-mortem dissections (Fernández-Miranda et al., 2008; Lawes et al., 2008).

A second conception comes from researchers who have reconstructed the longitudinal pathways with dMRI tractography based on anatomical priors from macaque tract-tracing experiments (Figure 3 right panel) (e.g., Barbeau et al., 2020; Frey et al., 2008; Makris et al., 2005; Warrington et al., 2020). Considering that neural tract-tracing is the gold standard for determining the precise origins and terminations, and axonal courses of brain connections, the macaque model that separates the peri-sylvian longitudinal fibres into the AF, SLF I, II and III bundles (i.e., Frey et al., 2014) would constitute the most anatomically accurate model of the primate longitudinal fibre system at present (Figure 3 right panel). Critically, it remains questionable how generalizable this macaque model is to the human brain, since after all language (which is associated with the AF) is not present in the macaque. One strategy, adopted by Petrides and colleagues (e.g., Frey et al., 2014; Petrides and Pandya, 1988; 2009), was to perform tract-tracing from macaque cortical regions whose human counterparts are known, based on cytoarchitectonics, which facilitates the macaque-human transfer of the anatomical findings. Indeed, by performing in-vivo dMRI tractography in the human brain using carefully defined anatomical landmarks that are known to be homologous to the macaque, recent researchers have been able to replicate the macaque AF, SLF I, II, III and MdLF fibre tracts in the human brain (e.g., Barbeau et al., 2020; Frey et al. 2008; Luo et al., 2020; Makris et al., 2005). Note that as defined by Luo et al., (2020) the MdLF “courses through the superior temporal gyrus (STG) and inferior parietal lobule (IPL) connecting STG (BA 22, 42) and temporal pole (TP, BA 38) principally with the angular gyrus (AG, BA 39). The MdLF is distinct from other long association fiber tracts connecting the frontal, parietal or temporal lobes such as the superior longitudinal fascicle II (SLF II) and SLF III or the arcuate fascicle (AF), respectively, and the extreme capsule (Makris et al., 2009)”.

How then might we reconcile those two legitimate conceptions of the AF? A potential hypothesis is that “the long direct segment”, “the anterior indirect segment”, and “the posterior indirect segments” defined in the first conception might correspond to “the AF”, “the SLF II and/or III”, and “posterior part of the AF and the posterior part of the middle longitudinal fasciculus (MdLF; based on Petrides, 2014; see also e.g., Latini et al., 2021) as defined in the second conception respectively. This could be verified via a virtual dissection experiment using dMRI tractography to test whether it is possible to obtain the macaque-derived model of the AF from Catani’s model (i.e., first conception) using additional region-of-interests, since currently, the latter has been derived using a two-ROI, or three-ROI approach to detect each of the three segments (e.g., Catani et al., 2005; Forkel et al., 2014).

Figure 3: Contemporary Models of the AF fibre pathways in humans.

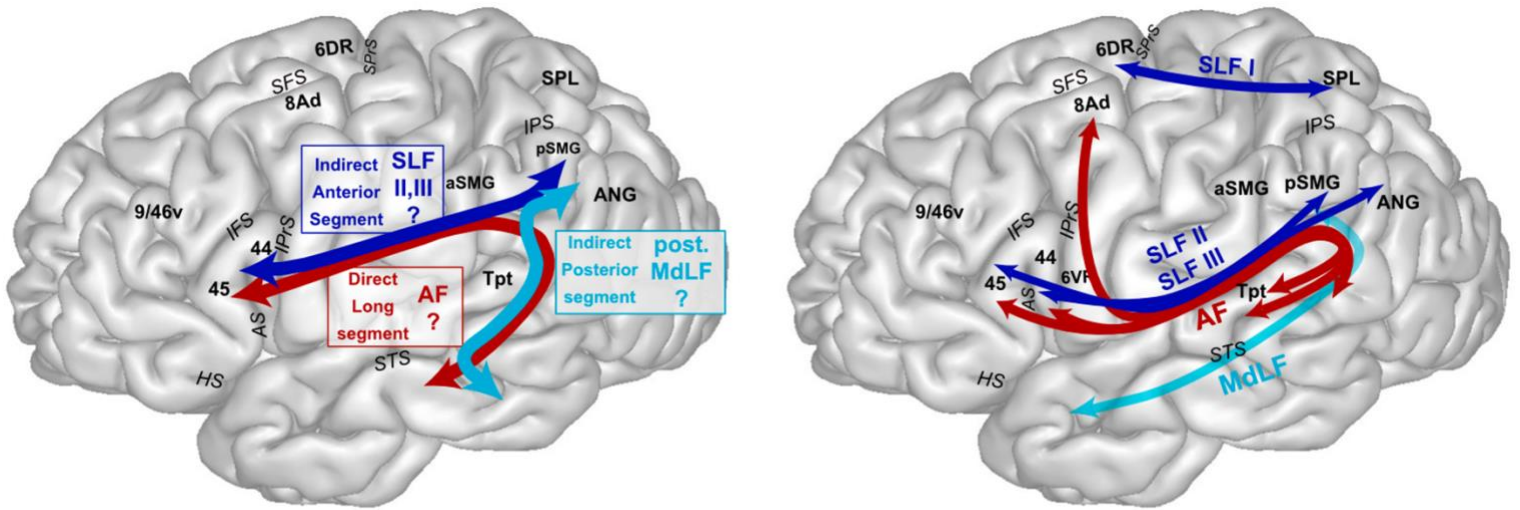


Figure 3: Contemporary Models of the AF fibre pathways in humans. Left: Dissection based human tractography AF/SLF model. Figure adapted from Catani et al., (2005). Right: Tract-tracing based human AF/SLF model. Figure adapted from Petrides (2014). Potential correspondences between figures are displayed.

SLF I, II, III= First, second and third branch of the Superior Longitudinal Fasciculus, AF= Arcuate Fasciculus; SFS=Superior Frontal Sulcus; SPrS=Superior Precentral Sulcus; IPrS=Inferior Precentral Sulcus; IFS=Inferior Frontal Sulcus; AS=Ascending branch of Sylvian Fissure; HS=Horizontal branch of Sylvian Fissure; IPS=Intra-Parietal Sulcus; Tpt=Temporal-parietal area (BA22); aSMG=anterior Supra-Marginal Gyrus (BA40); pSMG=posterior Supra-Marginal Gyrus (BA40); ANG=Angular Gyrus (BA39); SPL=Superior Parietal Lobule (BA7).

## Box 1: Partially overcoming diffusion MRI tractography limitations

To overcome diffusion MRI limitations, several methods have been adopted by researchers using dMRI tractography:

(1) *Increasing the resolution and sample sizes of in-vivo dMRI studies, or to include ex-vivo brains* that allow longer acquisition times and higher spatial resolutions (e.g., Barrett et al., 2020). This will reduce the rates of false positives, and increase reproducibility during tract reconstructions.

(2) *Focusing fibre reconstructions on the main bodies of the tracts (which are less ambiguous to detect) rather than their cortical projections* (e.g., Bryant et al., 2020; Warrington et al., 2020). This will enable a more stable and reproducible set of tracts to be reconstructed across individuals, and even across species, for comparative studies (Mars et al. 2019).

(3) *Relying on precise anatomical priors when defining tracts for reconstruction* (e.g., Barbeau et al., 2020; Catani et al., 2005). Here, the tracking of the streamlines would be constrained using prior anatomical knowledge inherited from ex-vivo techniques (i.e., defining seeds, targets and exclusion masks based on knowledge about the origin, terminations and course of a particular fibre tract), resulting in a more accurate “dissection” of specific fibre pathways using diffusion tractography. However, precise anatomical prior also cancels out variability. And, last but not least,

(4) *Advancement of tractography models* (e.g., Dell’Acqua and Tournier, 2019). New models are emerging, in order to overcome Diffusion tensor model limitations when reconstructing tract (such as crossing fibres). These models calculate multiple fibre orientations for tractography application (such as “spherical deconvolutions”).

## Part II: Four questions about the evolution of the Arcuate Fasciculus across primates

Current comparative work has revealed four key findings about the unique evolution of the human AF anatomy and compared these to macaques and chimpanzees. First, studies have proposed that the frontal cortical terminations of the AF have changed in the human brain (e.g., Schmahmann and Pandya, 2006). Second, studies have proposed that the temporal cortical terminations of the AF have expanded in the human brain (eg. Eichert et al., 2019). Third, studies have proposed a greater Dorsal (which includes the AF) *versus* Ventral Pathway expansion across primate evolution (e.g., Rilling et al., 2008). Fourth, studies have proposed an absence of left-lateralisation in the monkeys AF, that is present in humans (e.g., Rocchi et al., 2021). Here, we will re-examine the evidence for these findings with respect to their AF conceptions and discuss their implications for language evolution theories.

Figure 4: Illustration of the possible AF's frontal (Question 1) and temporal (Question 2) cortical terminations in the macaque brain

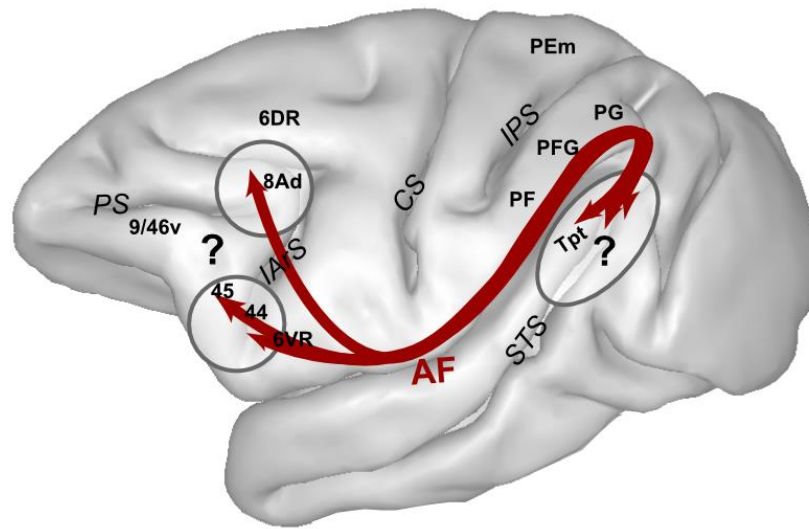


Figure 4: Schematic illustration of the homologue's main language pathways in the macaque monkey brain. Illustration of the course and terminations (arrowheads) of the Arcuate Fasciculus (AF) in red. Grey circles indicate the possible terminations as discussed in the literature (see table 1). Question marks highlight the research questions.

### Question 1: Are AF terminations in the inferior frontal cortex (Broca's area) present in the human-, but not in the nonhuman primate brain?

Classically, the human AF connects to the Broca's area in the inferior frontal cortex (Brodmann's areas 44 and 45). These frontal regions are significant for language because they are involved in the high-level control of orofacial and vocal motor responses that are needed for language production (Loh et al. 2020). Direct AF connections to these frontal regions thus provide access to the key motor actions (e.g., articulation, facial expressions, gestures) needed for language expression in humans (Petrides and Pandya, 2009). Whether the same AF frontal projections exist in nonhuman primates (NHPs) is therefore a crucial question that would shed light on why language exists only in humans. Currently, there are several theoretical accounts based on how a lack of AF frontal terminations in the nonhuman primate brain could explain the uniqueness of human language (e.g., Aboitiz and García, 1997; Friederici 2017; Pulvermueller 2018; Schmahmann and Pandya, 2006). However, the most recent data from tract-tracing experiments (e.g., Frey et al., 2014) appear to contradict these accounts as AF connections to macaque areas 44 and 45 were demonstrated. Here, we carefully review past and present tract-tracing studies of the macaque peri-sylvian longitudinal fibre system to resolve the question about whether AF inferior frontal terminations exist in the macaque brain.

Based on the landmark tract-tracing investigation in the macaque brain, Petrides and Pandya (1988) had traced AF terminations into the dorsal part of the frontal cortex (dorsal areas 6, 8A and prefrontal area 46), but not to the inferior frontal cortex, where Broca's area's homologue lies. Instead, the inferior frontal cortex only connected to the parietal cortex via the SLF II and III. This result remained important in the field over the next two decades: it was

1 included and reproduced in the highly influential atlas of fibre pathways of the macaque brain by  
2 Schmahmann and Pandya (2006; see Fig 13-5 on pp 407 for a depiction of the AF) and served as  
3 the anatomical reference for several dMRI studies (e.g., Thiebaut de Schotten et al. 2012) (Figure  
4 5 Left panel). Critically, this discovery had led to several theoretical accounts suggesting that the  
5 change in AF's frontal projections from dorsal part of the frontal cortex in nonhuman primates to  
6 Broca's area in humans could explain the emergence of human language (e.g., Aboitiz and García,  
7 1997; Friederici, 2017; Pulvermueller, 2018; Schmahmann and Pandya, 2006). For instance,  
8 Schmahmann and Pandya (2006) proposed that the macaque AF might not be a "language bundle"  
9 (pp 408) as it does not connect the homologues of Broca's (inferior frontal cortex) and Wernicke's  
10 areas (posterior superior temporal gyrus), but instead the caudal temporal area Tpt (which is  
11 associated with auditory-spatial processing) and the dorsal frontal areas (that is associated with  
12 oculomotor and attentional control). Due to the lack of a direct dorsal connection of Broca's  
13 homologue with the temporal lobe, but rather with the parietal lobe, Aboitiz and García (1997)  
14 proposed the parietal lobe to play a fundamental element in language evolution, linking indirectly  
15 the two key structures (Broca's and Wernicke's) by means of the SLF II and III. In their view, it  
16 "permitted the development of an interphase between the auditory processing device and a working  
17 memory circuit for complex vocalisations that includes inferoparietal and ventral premotor  
18 regions". Other theoretical accounts suggested that the absence or reduced degree of the AF's  
19 terminations in the inferior frontal cortex (area 44) in NHPs led to the lack of syntax (Friederici,  
20 2017) or action-perception representations (Pulvermueller, 2018) that are necessary for language.

21 A conceptual change in the views of the AF's frontal terminations happened when the  
22 cytoarchitectonical equivalence of Broca's area 44 was established in the macaque (Petrides et al.,  
23 2005), which prompted a series of tract-tracing experiments to reinvestigate the frontal-parietal-  
24 temporal pathways with new and more precise tracer injection sites (Petrides, 2014) (Figure 5).  
25 Following a detailed comparative analysis of the cytoarchitecture of macaque and human frontal  
26 cortex, Petrides and Pandya (Petrides et al. 2005a; b; Petrides and Pandya 1999; 2002) established  
27 critical correspondences between the various frontal areas between the two species. In particular,  
28 this work revealed for the first time the precise organisation of the homologues of Broca's area 44  
29 and 45 in the macaque brain. With these knowledge, the longitudinal fibre connections between  
30 the frontal, parietal and temporal lobe were re-investigated, leading to the major discovery of the  
31 existence of long arching fibres that directly connect macaque areas 44 and 45 to the caudal  
32 temporal cortex (Frey et al., 2014; Petrides and Pandya, 2009). Petrides explains in his 2014 book,  
33 that no terminations to area 44 and 45 were previously found, because the initial injection site was  
34 not placed ventrally enough within the temporal lobe (Petrides and Pandya, 2009; Petrides, 2014)  
35 (see Figure 5 right panel that shows each different injection site with an arrowhead and the  
36 resulting course with a different colour)

37 In fact, this work revealed three AF branches projecting to the frontal cortex (see Petrides  
38 2014, pp 160, Fig 52): 1) A dorsal branch that connects from the dorsal part of the superior  
39 temporal gyrus to dorsal frontal area 8Ad and area 6; 2) A second branch that connects from the  
40 ventral part of the superior temporal gyrus, and the upper bank of the superior temporal sulcus to  
41 area 44; and 3) A last branch that connects from the caudal superior temporal gyrus and the  
42 adjacent bank of the superior temporal sulcus to area 45 (Figure 5). This result was recently  
43 reproduced in comparative dMRI studies, in-vivo and ex-vivo, in macaques and chimpanzees  
44 (Balezeau et al. 2020; Barrett et al. 2020; Bryant et al. 2020; Eichert et al. 2019, 2020; Rilling et  
45 al. 2008; Roumazeilles et al. 2020) and on a functional level using effective connectivity (Rocchi  
46 et al., 2021).

As such, based on the above recent tract-tracing data, it is clear that AF connections to the inferior frontal cortex exist in the macaque brain, as in humans, suggesting that the so-called human language pathway between Broca's and Wernicke's areas might be more preserved in nonhuman primates than previously thought. Further supporting this conserved primate AF-inferior frontal cortical connectivity, recent studies have also reported common roles of macaque and human frontal areas 44 and 45 in cognitive vocal control (Loh et al., 2017; Loh et al., 2020). As such, contemporary theories that explain the uniqueness of human language based on the absence of inferior frontal projections of the AF must be revised accordingly. If AF frontal terminations are highly conserved, the question arises why language has evolved in humans alone? This important open question remains to be addressed by future research.

Figure 5: Knowledge progress about the macaque Arcuate Fasciculus from tract-tracing experiments.

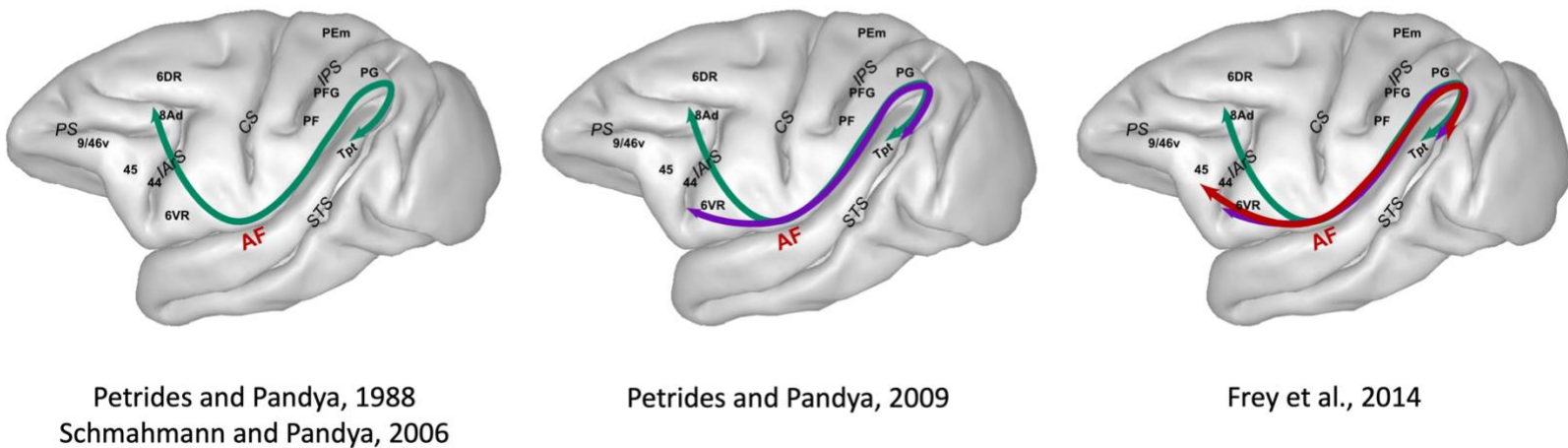


Figure 5. Knowledge progress about the macaque Arcuate Fasciculus courses and terminations from in tract-tracing experiments. (Left) Injections into the macaque caudal superior temporal gyrus (area Tpt) reveal the dorsal branch of the AF (Petrides and Pandya, 1988; Schmahmann and Pandya, 2006). (Middle) Injections into the ventral part of Tpt, and the upper bank of the STS reveals a second branch of the AF to area 44 (Petrides and Pandya, 2009; Case 5). (Right) More ventral injections into the caudal superior temporal gyrus and STS reveals a third branch of the AF to area 45 (Frey et al., 2014). AF=Arcuate Fasciculus; IArS=Inferior Arcuate Sulcus, PS=Principal Sulcus; CS=Central Sulcus; IPS=Intra-Parietal Suclus; STS=Superior Temporal Sulcus. PF; PEm; PFG; PG are cytoarchitectonic labels for the parietal areas based on Von Economo and Koskinas (1925)

## Question 2: Do human AF terminations extend further into the temporal lobe than in nonhuman primates?

In the human brain, the AF arches around the Sylvian fissure, and extends beyond the posterior part of the superior temporal gyrus, where the Planum Temporale and Tpt lies, and stretches into the middle temporal gyrus (e.g., Rilling et al., 2008; Catani et al., 2005; see Figure



3 left panel), and anterior temporal areas (e.g., Eichert et al., 2019). This middle part of the temporal cortex is associated with semantic verbal (or multimodal) information (Binder et al., 2009), and as such, this implies that the human AF provides the frontal language areas access to semantic information in the middle temporal gyrus. The difference in temporal terminations between human and NHP is therefore crucial, in order to investigate potential evolutionary divergences, which, in turn, influence functional theories about language evolution. It is, however, not to be excluded that this human-unique feature could be: 1) due to the expansion of the temporal lobe alone (Eichert et al. 2019; Van Essen and Dierker 2007); or 2) due to methodological differences (Latini et al., 2021; Petrides, 2014) (Figure 4).

In their diffusion MRI work, Rilling and colleagues (2008; 2012) described major changes during primate evolution with an increased temporal lobe projection in the human brain and related this finding to language evolution. While the human AF reaches the MTG, the macaque AF terminates at the caudal end of temporal area 22, near the temporo-parietal junction. As such, Rilling et al. (2008) suggested that the human AF is unique in its extended projections into the middle temporal gyrus. This finding has become the basis of a number of models of human language processing. For example, in their model for “Social Communication And Language Evolution and Development” (SCALED), Catani and Bambini (2014) linked the lack of the AF’s deep terminations into the middle temporal gyrus and anterior temporal areas (long temporal tail) in NHPs to the absence of syntax processing and reduced semantics processing.

At the same time as Rilling et al. (2008), Van Essen and Dieker (2007) also proposed a theory of primate brain evolution that described a cortical expansion across primates using certain key regions of interest as common landmarks. They proposed that if the macaque brain was inflated to the size of a human brain, some areas would be too small in comparison to their human counterparts. Thus, these areas, including the middle temporal cortex, must have expanded disproportionately in size during evolution to match the actual human brain organisation. As such, this temporal cortical surface expansion could explain the apparent expansion of AF terminations into the middle part of the temporal lobe in the human brain.

By examining the surface projections of the temporal association tracts across primate species, a series of studies from Rogier Mars’ laboratory tackled the question of whether the observed extension of the AF’s termination into the human temporal lobe was due to cortical expansion alone, or also to an actual extension of the tract into new cortical regions (Bryant et al.; 2020; Eichert et al., 2019; 2020; Mars et al., 2021; Roumazeilles et al., 2020; Warrington et al., 2020). Eichert et al., (2019) first extracted the main body of the AF tract in both the macaque and human brain using common tractography protocols (Warrington et al, 2020). The cortical surface projections from these AF tract bodies were then computed in each species. Next, similar to Van Essen and Dieker (2007), interspecies cortical surface transformations were computed using species-common cortical landmarks. Finally, by applying these cross-species surface transformations on the macaque AF tract projection maps and comparing them to the actual human AF maps, they examined whether cortical surface expansion alone would predict the apparent expansion of AF tract projections from macaque to humans (see also Eichert et al., (2020) for a similar method based on T1w/T2w maps alignments). This way, they demonstrated that cortical surface expansion alone cannot account for the expansion of the AF’s tract projections in the human temporal lobe. (Eichert et al. 2019; 2020). The authors therefore conclude that the human AF underwent additional changes in its course by invading new middle and anterior temporal areas. Providing a full picture of how the AF and temporal cortex had evolved across primates, the same group of researchers proposed that the middle temporal region first underwent a significant

1 cortical expansion in the great apes lineage, and subsequently, in human evolution, the AF extends  
2 into these new temporal territories, resulting in the expanded cognitive/language abilities in  
3 humans (Roumazeilles et al., 2020)

4  
5 Echoing these findings, Barrett et al. (2020) found no terminations exceeding the Tpt area  
6 to connect with frontal areas in macaque *ex-vivo* dMRI data, which were present in humans. They  
7 state that the AF shows the most striking differences among the various association tracts between  
8 humans and monkeys, with monkeys sharing only a small subcomponent of the AF (roughly the  
9 body of the tract) with humans. As such, consensus lies in the fact that the monkey's AF's temporal  
10 tail terminates in the Tpt/Planum Temporale area (e.g., Schmahmann and Pandya, 2006, but see  
11 also Schmahmann and Pandya, 2007). This morphology is incomparable with that of the human,  
12 where it exceeds the middle temporal gyrus and anterior temporal areas.

13 However, very recent studies found for the first time connections between the functionally  
14 defined auditory cortex, and Broca's homologue (Areas 44, 45). This result highlights a slightly  
15 more anterior temporal projection of the AF than previously found, since the auditory cortex is  
16 located in the middle and posterior superior temporal lobe, anterior to the Tpt/Planum Temporale  
17 area (where the AF has been shown to terminate in previous works). Based on this direct  
18 connection between the auditory cortex and Broca's homologue, the authors speculate "that this  
19 dorsal auditory pathway is involved in not just spatial processing in the classical sense but also  
20 sound and vocal patterning in the time domain", a function that had previously been exclusively  
21 linked to the Ventral Pathway (see Question 3). Note that in this description, the human AF also  
22 terminates in the posterior superior temporal lobe and does not exceed into the middle- or anterior  
23 parts of the temporal lobe.

24 However, the consensus that the human AF arches into the middle, anterior and inferior  
25 parts of the temporal lobe is questioned by some authors (Latini et al., 2021; Maldonado et al.,  
26 2013; Petrides 2014; Wang et al., 2013). In fact, in the monkey brain tract tracer studies have  
27 demonstrated that monosynaptic AF connections terminate around the Tpt/Planum Temporale  
28 area. Fibres arching further ventrally in the temporal lobe (i.e., temporo-parietal fibres like the  
29 posterior middle longitudinal fasciculus (MdLF), see Part I "Late 20th Century: Neural tract-  
30 tracing...") are independent and build synapses with the AF (Distler et al. 1993; Petrides 2014;  
31 Seltzer and Pandya 1984). The possibility cannot be excluded, that this morphology may be similar  
32 in the human brain, with the human AF also originating in area Tpt/Planum Temporale if defined  
33 monosynaptically. However, with diffusion MRI, one is unable to distinguish between a  
34 monosynaptical AF terminating in area Tpt/Planum Temporale or a polysynaptical AF  
35 intermingled with tracts running through the temporal lobe, like the posterior MdLF (as proposed  
36 by Latini et al., 2021; Maldonado et al., 2013; Petrides 2014; Wang et al., 2013). In other words,  
37 in this view, the human AF might be losing its predominant "arch" (which might correspond to  
38 the posterior indirect segment (Catani et al., 2005)) due to methodological issues (Figure 3). It is  
39 possible that this hypothesis influenced the most recent comparative investigations (Balzeau et al.,  
40 2020; Rocchi et al., 2021). Indeed, in these recent studies, seeds for the human AF are placed in a  
41 way that the AF is terminating more posteriorly and less ventrally in the temporal lobe than in  
42 previous work (Barrett et al., 2020; Eichert et al., 2019; Rilling et al., 2008; 2012), but terminating  
43 like their monkey homologue in the auditory cortex (Balzeau et al., 2020; Rocchi et al., 2021).

44 If we consider the hypothesis that the temporal projections of the human AF indeed  
45 terminate at the posterior superior temporal cortex as in the macaque (e.g. Latini et al., 2021;  
46 Maldonado et al., 2013; Petrides, 2014; Wang et al., 2013), what, then, might be the cause of the



1 apparent differences in temporal lobe tractography between humans and monkeys? One possibility  
2 could be that, besides the AF, the various tracts that run through the temporal lobe (e.g., the Inferior  
3 Longitudinal Fasciculus (ILF) Inferior Fronto-Occipital fasciculus (IFOF), the Middle  
4 Longitudinal Fasciculus (MdLF), Uncinate Fasciculus (UF), Extreme Capsule Fasciculus (ECF)  
5 etc.) might have expanded along with human temporal lobe evolution (Braunsdorf 2021). These  
6 tracts are considered part of the Ventral pathway and might be interconnected with the AF  
7 (Braunsdorf 2021; but see Rilling et al., 2012, for the exclusion of potential ILF contamination  
8 during the reconstruction of the AF). In other words, this idea is in line with a human-specific  
9 evolution of temporal lobe connectivity that is not, however, due to the AF, but to other tracts.

10 In summary, current consensus lies in the divergence of AF temporal terminations.  
11 Monkeys possess terminations in the most posterior part of the temporal lobe, even though recent  
12 advances pushed these terminations more anteriorly into the auditory cortex (Balezeau et al.,  
13 2020). In humans, the AF extends into the middle part of the temporal cortex, and for some authors  
14 more anterior-inferiorly into anterior temporal areas. Even though the possibility has to be  
15 highlighted that this extension, relative to macaques, might be due to polysynaptic connections  
16 with other Ventral Pathway fasciculi such as the posterior part of the MdLF. In the future,  
17 investigations have to show whether this difference in temporal terminations between human and  
18 NHP is genuinely crucial and accounts for the evolution of language in the human lineage.

### 22 **Question 3: Has the Dorsal Pathway, relative to the Ventral Pathway, expanded** 23 **significantly in humans?**

24  
25 Recent language models depict a Dual Pathway model (Hickok and Poeppel, 2004) which  
26 emulates the well-known visual system division into a Ventral and a Dorsal Pathway. In this  
27 model, both pathways connect the inferior frontal to the temporal lobe, with the Dorsal Pathway  
28 curving dorsally around the Sylvian Fissure, and Ventral Pathway passing ventral to the Sylvian  
29 fissure. Traditionally, at the functional level, the Dorsal Pathway is involved in mapping sound to  
30 articulation, while the Ventral Pathway is involved in mapping sound to meaning (Hickok and  
31 Poeppel, 2004; Saur et al., 2008). There has been great interest recently in understanding how these  
32 pathways have differentially evolved across species due to their direct relevance to human  
33 language functions. Due to limitations in disentangling fibre bundles in dMRI tractography, it is  
34 anatomically unclear exactly which fibre bundles form these two broad conceptual pathways.  
35 While it is generally accepted that the AF together with SLF II and III make up the Dorsal language  
36 Pathway (see part I), the exact tracts that constitute the Ventral Pathway remain ill-defined in the  
37 literature. As such, many diffusion MRI studies thus cannot account for a clear separation of the  
38 individual tracts constituting the Dorsal and Ventral Pathways and have therefore instead based  
39 their interpretations on this broad Dorsal/Ventral dichotomy (e.g., Balezeau et al., 2020; Eichert et  
40 al., 2019; Rilling et al., 2008; 2012). For this reason, we also keep here the appellation “Dorsal  
41 and Ventral pathways” as in the original articles (e.g., Rilling et al., 2008; 2012). While both  
42 pathways connect similar areas, it is the Dorsal Pathway that is dominant in humans (i.e., stronger  
43 structural connectivity than the Ventral Pathway) (Rilling et al., 2008). A crucial question is  
44 therefore whether the same organisation is shared with monkeys.

45 Currently, there are several theoretical accounts based on how a reduced ~~or-inversed~~  
46 Dorsal- *versus* Ventral dominance in nonhuman primates could explain the uniqueness of human

1 language (e.g., Friederici, 2017; Pulvermueller, 2018; Rilling et al., 2008; 2012). However, with  
2 differences in seeding and the addition of functional connectivity, recent investigations have  
3 challenged this view and show increased connectivity on the structural side for the Dorsal Pathway  
4 and an equal dominance of both pathways on the functional side in monkeys (Balezeau et al., 2020;  
5 Rocchi et al., 2021). Here, we review tractography and effective connectivity studies of the Dorsal  
6 and Ventral Pathways to investigate whether or not a stronger ventral-than-dorsal ratio exists in  
7 the nonhuman primate's brain.

8 In their pioneering work, Rilling and colleagues (2008) compared the Dorsal and Ventral  
9 Pathway across primate species, using dMRI and a probabilistic multi-fibre diffusion model. They  
10 found a stronger (more streamlines) Dorsal-than Ventral Pathway in human, which was not the  
11 case in macaques and chimpanzees. In fact, for macaques a very weak (i.e., low number of  
12 streamlines) Dorsal Pathway was described, where frontal terminations were found in area 46 and  
13 delicately in area 44 (part of Broca's homologue). Temporal terminations were described only in  
14 the most posterior end of the temporal lobe in the junction with the parietal lobe. In contrast, a  
15 strong Ventral Pathway was highlighted with terminations in area 44 and 45 (Broca's homologue)  
16 as well as in the frontal operculum and across the temporal lobe (see Figure 4, left panel). In  
17 humans the authors described an opposite organisation with a Dorsal Pathway stronger than the  
18 Ventral one, connecting both similar areas (Rilling et al., 2008). In addition, in comparisons to  
19 chimpanzees, they described qualitatively stronger connections to Broca's area (area 44 and 45)  
20 via the Dorsal Pathway, than via the Ventral pathway (Rilling et al., 2012) in humans. The authors  
21 concluded that the Dorsal pathway changed significantly more than the Ventral Pathway in human  
22 evolution, due to the overall strengthening, greater strength than the Ventral Pathway, stronger  
23 area 44 and new area 45 terminations as well as extended terminations into (and beyond) the  
24 middle part of the temporal lobe (replicated e.g., by Eichert et al., 2019) (*Figure 6 left panel*). The  
25 authors further conclude that the proportional strengthening of the Dorsal Pathway might have  
26 contributed to transfer lexical-semantic information and some elements of syntax in the evolution  
27 of human language. In contrasts, its monkey's homologue would be rather involved in low-level  
28 spatial processing of sounds (Aboitiz, 2018; Rilling et al., 2008), similar to the view of  
29 Schmahmann and Pandya (2006) (Question 1 above). Also, the great extent of temporal lobe  
30 information like auditory information would be exclusively conveyed via the Ventral Pathway in  
31 monkeys (in agreement with functional and tract tracing results (Romanski et al. 1999)). This  
32 finding and interpretation had a conceptual impact on several theories trying to retrace the  
33 evolution of human language. Here we will focus on two of them (Friederici, 2017; Pulvermueller,  
34 2018).

35 A first prominent theory comes from Angela Friederici as highlighted in her 2017 book  
36 "Language in the brain" (Friederici, 2017). She proposes that complex syntax and therefore human  
37 language developed in human only, due to the strengthening of the Dorsal Pathway with extension  
38 of terminations into area 44 (part of Broca's area). The reasoning is the following: The Dorsal  
39 Pathway conveys complex syntax in the human brain due to the connection to area 44 (Friederici  
40 et al. 2006), whereas the Ventral Pathway conveys semantic aspects of language and simple syntax  
41 due to the connection to area 45 and the frontal operculum (Friederici, 2017). Monkeys however  
42 show non or only weak dorsal connections to area 44 (Rilling et al., 2008), but have ventral  
43 termination to this area. Therefore, they don't show complex syntax, but aspects of semantics and  
44 simple syntax (Friederici, 2017). Indeed, behavioural data indicates that monkeys fail to show the  
45 processing of complex syntax, but showed aspects of semantics (Friederici, 2017). In addition,  
46 behavioural data in developmental studies highlight that infants only acquire complex syntax by

1 the age of seven, exactly when the most anterior portion of the Dorsal Pathway matures and  
2 connects to Broca's area (Brauer et al. 2013; Brauer et al. 2011; Friederici 2017; Perani et al.  
3 2011). In contrast, the Ventral Pathway connection into this area is already present at that age  
4 (Brauer et al. 2013; Brauer et al. 2011; Friederici 2017; Perani et al. 2011).

5 However, there are also some limitations to this theory:

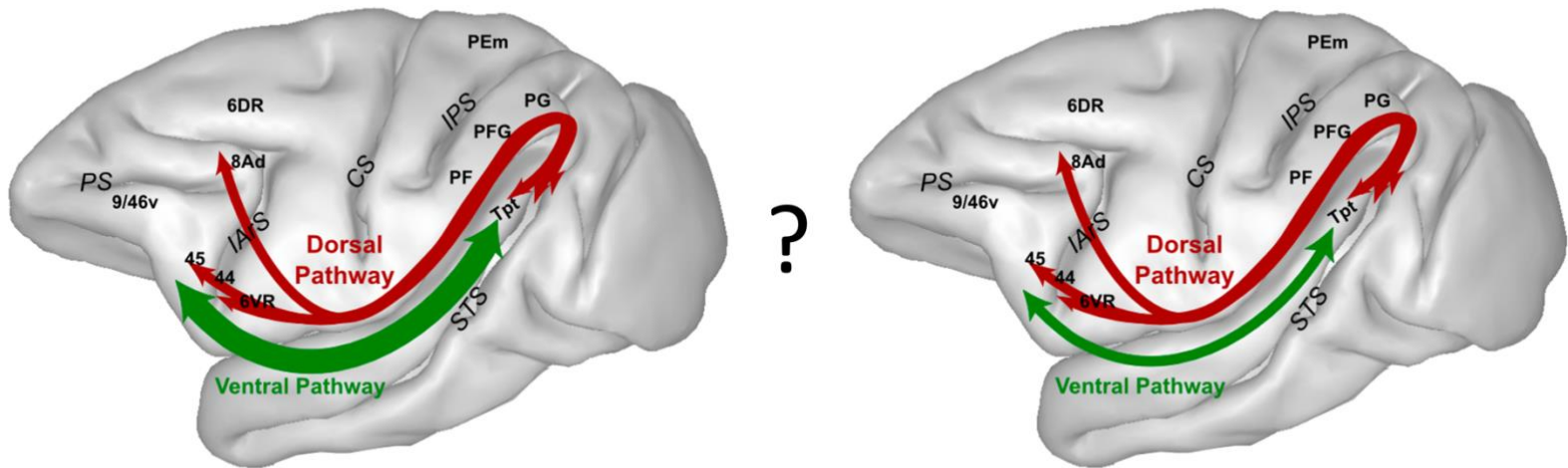
6 First, several dMRI studies interpret the developmental data differently (Dubois et al. 2009, 2016).  
7 In their view Dorsal Pathway's terminations to Broca's area are present before the age of 7, but  
8 intractable in dMRI due to their immature microstructure. Second, more recent behavioural studies  
9 highlight that monkey can "break to the syntax barrier" (Ferrigno et al. 2020; Jiang et al. 2018;  
10 Malassis et al., 2020; Fitch 2018).

11 A second recent and influential theory based on a weaker Dorsal Pathway in NHP was  
12 proposed by Pulvermüller (2018). He hypothesised that the overall strengthening, and the  
13 evolutionary new and stronger area 44 termination of the Dorsal Pathway enabled frontotemporal  
14 sensorimotor shortcuts, connecting further than just next neighbouring areas. This way, humans  
15 can better bind action and perception into circuits that support cognitive functions important for  
16 language processing, like working memory, prediction, attention, and combination, including  
17 abstract rule formation. In this work, action representations lie at the heart of language processing.

18 Very recently however, changes in techniques and seeds for tracking have provided the  
19 first arguments against ~~have-challenged~~ the view of a stronger-Ventral-than-Dorsal-Pathway in  
20 monkeys. In a diffusion MRI studies across primate species, Balezeau and colleagues (2020) used  
21 a functionally defined auditory cortex as the seed for tractography. This way, they highlighted a  
22 Dorsal Pathway connection into the auditory cortex, which was until then reserved to the Ventral  
23 Pathway (Rilling et al., 2008; Romanski et al, 1999). The authors therefore suggest that the monkey  
24 AF is not solely involved in low-level processes but instead important for sound and vocal  
25 patterning in the time domain (Balezeau et al., 2020).

26 Furthermore, on the functional level, using functional MRI while electrically stimulating  
27 the same auditory cortex (termed "effective connectivity"), Rocchi and colleagues (2021) reported  
28 that Dorsal and Ventral Pathways connect to the same degree to Broca's homologue (area 44 and  
29 45) and the frontal operculum. In other words, even though the Ventral Pathway was shown  
30 stronger on a structural level (Rilling et al., 2008; 2012), functionally both pathways have the same  
31 strength. The authors suggest therefore that language abilities allowing humans to name,  
32 conceptualize and thus better remember sound (Schulze et al., 2012) would be shared across  
33 primates (Figure 6 right panel). Based on the above recent data, the long-standing bulwark of a  
34 reduced Dorsal Pathway dominance that could explain language evolution is beginning to crackle.  
35 As a result, contemporary theories that explain the uniqueness of human language based on the  
36 ~~inverted~~ proportional strength of the Dorsal and Ventral Pathway, might benefit from a revision  
37 ~~be revised accordingly~~. While structural differences in strength may persist between the human  
38 and the monkey Dorsal Pathway, this may not be the case functionally. A question that exceeds  
39 the current review is to investigate whether structural differences are therefore of importance on  
40 other functional levels as investigated by Rocchi et al. (2021).

1 Figure 6: Dorsal- versus Ventral Pathway strength (question 3)



2 Figure 6: Dorsal- versus Ventral Pathway strength (question 3). (Left) A dominance in strength  
 3 (in structure and function) in the Ventral Pathway (green) against the Dorsal Pathway (red) (e.g.,  
 4 Rilling et al., 2008; 2012). (Right) Equal strength (in structure and function) between both  
 5 pathways (e.g., Rocchi et al., 2021).

6

7 **Question 4: A continuous evolution of the AF across primates - Could AF lateralisation be**  
 8 **the key to human language?**  
 9

10 In humans, the AF is strongly lateralised towards the left hemisphere (e.g., Thiebaut de  
 11 Schotten et al., 2011). This structural lateralisation is of interest, because the majority of the  
 12 language system is also functionally left-lateralised in the brain (e.g., Toga and Thompson 2003).  
 13 Whether such a same lateralisation exists, or not in nonhuman primates is therefore a crucial  
 14 question that would shed light on the origins of human language lateralisation. Especially, since  
 15 the most recent investigations about the AF's anatomy across primates agree on a highly  
 16 evolutionarily conserved anatomy (see sections above). Thus, the question about anatomical  
 17 differences across primates, that might explain the human uniqueness for language, persists.  
 18 Authors of recent investigations highlight that the left AF lateralisation is the crux of the human-  
 19 specific distinction (Balezeau et al., 2020; Eichert et al., 2019; Rocchi et al., 2021). While carefully  
 20 reviewing the recent comparative brain lateralisation data, we conclude that this compelling idea  
 21 lacks sufficient sample size.

22 From early human development on, inter-hemispheric asymmetry was documented for  
 23 most language-related regions at both structural and functional level. In fact, language tasks elicit  
 24 greater activation in the left- than in the right hemisphere, even in newborns (e.g., Dehaene-  
 25 Lambert et al. 2002). Structural markers of such language brain lateralisation were proposed, like  
 26 the leftward lateralisation of the Planum Temporale, Broca's area or the Insula (e.g., Geschwind  
 27 and Levitsky 1968; Keller et al. 2010). However, the human uniqueness of such results was  
 28 challenged by finding some of these structural markers not only in apes (e.g., Hopkins et al. 2015)  
 29 but also in an Old World monkey, the baboon, in both adults and newborns (Becker et al. 2021a;

c; d; Marie et al. 2018). Regarding white matter in humans, volume of the SLF II and SLF III was shown larger in the right hemisphere (e.g., Amemiya et al. 2021; Thiebaut de Schotten et al. 2011), which is a phylogenetically shared feature with chimpanzees (Hecht et al. 2015). In contrast, the human AF (as defined by the “long direct segment”) was shown larger in the left hemisphere for a number of macroscopic and microscopic measurements like the number of streamlines, volume of the tract, fibre density and mean fractional anisotropy in adults and infants (Catani et al. 2007; Dubois et al. 2009; Lebel and Beaulieu, 2009; Nucifora et al. 2005; Powell et al. 2006; Thiebaut de Schotten et al. 2011; Takaya et al. 2015; Thiebaut de Schotten et al. 2011). Therefore, the structural AF asymmetry, and specifically its temporal terminations, is indeed discussed as a marker for the functional lateralisation of the language network (e.g., Catani et al., 2007; Powell et al., 2006; Takaya et al., 2015; but see also e.g., Silva and Citterio 2017; Vernooij et al. 2007) and was further supported by resting state MRI data showing functional lateralisation of the language network (Zhu et al. 2014). However, Verhelst et al. (2021) recently reported that the AF is left-lateralised, regardless of left or rightwards language lateralisation. Note that this lateralisation might be dependent on the technique used in tractography.

But what is known about the nonhuman primate’s AF lateralisation? Initially, Rilling et al. (2008) did not find any AF asymmetry in three macaque and four chimpanzee subjects. A follow-up study in chimpanzees including a higher number of subjects (Rilling et al., 2012), showed a left-lateralised AF for 26 subjects. However, this lateralisation in chimpanzee was weaker than in humans. Eichert et al., (2019) confirmed an absence of left asymmetry for the AF/Dorsal Pathway in four *ex-vivo* macaque with high resolution and one *in-vivo* macaque image. In contrast, a strong left lateralisation was reported for the Ventral Pathway, which was found slightly right lateralised in humans (Eichert et al., 2019). Similarly, Balezeau et al., (2020) reported no lateralisation for the Dorsal Pathway in three macaques but also no lateralisation for the three chimpanzee subjects. This latter finding was confirmed on a functional level by Rocchi et al. (2021) for human and macaque subjects. Despite only treating two macaque subjects, a phylogenetical discontinuity in lateralisation was highlighted, with humans being leftwards lateralised and monkeys showing no lateralisation. As one can easily see, studies that investigated lateralisation produced inconsistent results and are limited to small sample sizes. Indeed, it is common in neuroscience to use not more than two macaque individuals in a given study, which makes interpretation of a populational level lateralisation impossible. Tract tracing methods, for example, are not able to give a response to the questions, while dMRI tractography may do. A small sample-size also limits homogeneous groups in regards to handedness, which was shown to be linked to brain lateralisation (e.g., Hopkins et al., 2015; Meguerditchian et al., 2013), and which might influence therefore influence a population-level lateralisation. In addition, while lateralisation of homologous perisylvian language areas were found present in several ape species (i.e., Hopkins et al., 2015), these asymmetries were not detected in macaques (Lyn et al., 2011), which are commonly representing the family of Old world monkeys. In contrast, recent studies demonstrated that another Old world monkey, the baboon, possesses homologous asymmetries (Becker et al., 2021a;b;c;d; Marie et al., 2018). It remains unclear why cerebral asymmetries are commonly detected in baboons but not in macaques. Based on the above tractography data, the question about evolutionary continuities or discontinuities in AF lateralisation will need further investigations with 1) ample sample sizes also in regard to handedness and 2) the inclusion of the baboon as a representative of the Old world monkey family. Therefore, we predict that future studies will focus on this crucial question in language brain evolution.

### Part III: Discussion and Conclusion

The apparent literature about the evolution of the AF has been rapidly gaining in interest in very recent years and is used as a base for theories about language evolution. However, no clear consensus was found, due to differences in appellations and divergent anatomical courses and terminations, which are resulting in diverging definitions of what exactly the AF is. For example, anatomically, AF and SLF generally refer to the same tract. This is important as the label “AF” implies “language pathway” more than the labels SLF, MdLF or Dorsal Pathway. But note that all terms are used interchangeably throughout the literature. However, differences in terminology matters, because a same result could have different theoretical implications depending on its labelling and therefore its functional connotation. To illustrate this point we can do a thought experiment: If, during primate evolution, an increase of connectivity strength between the superior temporal lobe (Wernicke’s area) and the inferior parietal lobe (Geschwind’s region) (as in e.g., Aboitiz and García, 1997; Catani and Bambini, 2014) was documented, different functional interpretations could be drawn from this, depending on terminology. We could label this segment for example AF, SLF, MdLF or Dorsal Pathway. However, the functional implication would be the most associated with language evolution for the label “AF”, even though the segment is the same.

In this review, we have therefore carefully highlighted the history of AF anatomy across primate species and defined a clear current anatomical model. The AF/SLF complex can be divided into four distinct fibre bundles including: 1) the AF, which comprise in itself three fibre branches that arch dorsally around the Sylvian fissure and connect the posterior temporal cortex (area Tpt, posterior auditory cortex) with the frontal cortex (area 44, 45, area 8Ad and 6) and 2) the SLF I, II and III, which comprise the three longitudinal fibre bundles that interconnect distinct parts of the frontal and parietal cortex and are not arching around the Sylvian fissure. In addition, we focused on four discontinuous key changes in AF anatomy that are discussed to differentiate the human from the nonhuman primate. These are grouped in research questions concerning: frontal terminations (1), temporal terminations (2), the proportion of the Dorsal- versus Ventral Pathway strength (3) and lateralisation (4). These four key questions are of high importance, because their results in (dis)continuity influence theories about language evolution. Based on the latest results of these four key questions, we argue that the evolution of the AF anatomy is more continuous across primate species than previously described. When summarizing the four questions (see also Table 1), we can notice the importance of methodological changes that are pivotal in the conceptualisation of the AF anatomy:

- Historically, knowledge about the AF’s terminations (Question 1) relies strongly on invasive tract tracing studies in the monkey brain. Findings about differences in frontal termination in the macaque brain (no Broca’s terminations) compared to the human brain (Broca’s termination) led to theories about the specific frontal development of the human AF, important for language processing. However, human-like terminations into Broca’s homologue in monkeys were found when cytoarchitectonic equivalence was demonstrated and tract tracing injection sites changed. Therefore, the frontal termination’s organisation seems to be conserved during primate evolution, even though terminations might have become overall stronger in humans.
- The same applies to the temporal terminations (Question 2), which were initially found to be very extended in humans, exceeding the middle part of the temporal lobe into its anterior and inferior parts. New data from methodological changes indicate however, that first, the

1 monkey's terminations are also slightly more extended (tracking from the functionally  
2 characterised auditory cortex) and second, the human extension could be a technical  
3 artefact (possible entanglement with fibres running through the temporal lobe like the  
4 MdLF). Therefore, it is not excluded that the temporal termination's organisation might  
5 also be more conserved during primate evolution, even though the temporal connections  
6 might have become overall stronger in humans.

- 7 • Finally, pivotal changes in the question of the proportion of the Ventral- versus the Dorsal  
8 Pathway's strength (Question 3), can also be illustrated by changes in techniques (effective  
9 connectivity) and seed masks (tracking from the functionally characterised auditory  
10 cortex). Initially found to be very weak and not terminating into the auditory cortex, the  
11 Dorsal Pathway in monkeys might now ~~demonstrated~~ to be regarded as functionally as  
12 strong as the Ventral Pathway and also connecting with the auditory cortex. Therefore, the  
13 possibility is not excluded that the Ventral-versus-Dorsal Pathway strength organisation  
14 might be conserved during primate evolution, even though it is possible that the Dorsal  
15 Pathway might have become overall structurally stronger in humans.
- 16 • Using this method-based framework on the question of lateralisation (Question 4), we can  
17 say that currently only the progress of tractography methods and dMRI acquisition  
18 techniques could advance this question as tract-tracing methods are not adapted. In  
19 particular, a forthcoming development of tractography applied to a large-scale cohort of  
20 non-human primate data could further advance the question as it was already conducted in  
21 chimpanzees concerning the lateralisation of other tracts like the SLF (Hecht et al., 2015),  
22 or structural grey matter asymmetries (e.g., Hopkins et al., 2015). Therefore, the question  
23 of AF lateralisation across primates needs a specific attention in the future. Given existing  
24 results about the left-lateralisations of perisylvian regions in monkeys (Becker et al., 2021a;  
25 Marie et al., 2018), it is not excluded that a human-like left lateralisation could also be  
26 demonstrated in monkeys in the future, even though lateralisation might have become  
27 stronger in human evolution.

28 Last but not least, we do not want to understate the importance of comparative AF studies for  
29 the investigation of language brain evolution. In fact, the reviewed data points towards a  
30 progressive evolution of the AF during primate evolution, rather than great leaps that would  
31 indicate categorical changes of the AF's morphology between contemporary primate species. This  
32 progressive evolution could be expressed by a progressive 1) strengthening of connectivity, 2)  
33 expansion of terminations and 3) lateralisation of features that have been already present in the  
34 common ancestor of Old world monkeys, chimpanzees and humans. However, future studies must  
35 also focus on other tracts that might have undergone substantial changes during primate evolution  
36 and that are linked to language processing in humans. Possible tracts of interest are tracts that run  
37 through the temporal lobe (and might be constituents of the Ventral Pathway, depending on its  
38 definition), like the Inferior Longitudinal Fasciculus (ILF), Inferior Fronto-Occipital fasciculus  
39 (IFOF), the Middle Longitudinal Fasciculus (MdLF), Uncinate Fasciculus (UF), Extreme Capsule  
40 Fasciculus (ECF). But also, the Frontal Aslant, which connects the supplementary motor areas, the  
41 lateral superior frontal gyrus and the cingulate cortex to the inferior frontal gyrus could be a  
42 candidate to focus on (for a review: La Corte et al., 2021).

43 In conclusion, recent works in humans and nonhuman primates suggest that the AF course  
44 and terminations are more similar between species than described before. The remarkable human  
45 ability of language might therefore have evolved progressively in primate evolution, exactly as its  
46 underlying anatomic implementation in the brain. Or to put it in the words of Charles Darwin:

1 "Nevertheless the difference in mind between man and the higher animals, great as it is, certainly  
2 is one of degree and not of kind." (page 105: Darwin, 1871/1981). Future theories about language  
3 brain evolution should account for this continuous, and not discrete AF evolution.  
4

	<b>Tract tracer studies</b>		<b>Diffusion MRI</b>	
	<i>Reported</i>	<i>Not reported</i>	<i>Reported</i>	<i>Not reported</i>
<b>Broca's homologue terminations (monkey)</b>	Petrides and Pandya, 1988 Schmahmann and Pandya, 2006 Thiebaut de Schotten et al., 2012	Petrides and Pandya, 2009 Frey et al., 2014 Petrides et al., 2014	/	Rilling et al., 2008 Eichert et al., 2019 ; 2020 Barrett et al., 2020 Balezeau et al., 2020 Bryant et al., 2020 Roumazeilles et al., 2020
<b>Auditory cortex terminations (temporal lobe) (monkey)</b>	Schmahmann and Pandya, 2006 Petrides and Pandya, 2009 Frey et al., 2014 Petrides et al., 2014	/	Eichert et al., 2019 ; 2020 Barrett et al., 2020 Roumazeilles et al., 2020 Rilling et al., 2008 ; 2012	Balezeau et al., 2020 Rocchi et al., 2021
<b>Extended temporal lobe terminations (human)</b>	/	/	Maldonado et al., 2013; Wang et al., 2013; Petrides, 2014; Balezeau et al., 2020 Barbeau et al. ; 2020 Latini et al., 2021	Catani et al., 2005 Rilling et al., 2008 Thiebaut de Schotten et al., 2012 Eichert et al., 2019
<b>Ventral &gt; Dorsal Pathway (monkey)</b>	/	/	Rocchi et al., 2021	Rilling et al., 2008 ; 2012
<b>Lateralisation (chimpanzee)</b>	/	/	Balezeau et al., 2020	Rilling et al., 2008 ; 2012
<b>Lateralisation (monkey)</b>	/	/	Rilling et al., 2008 ; 2012 Eichert et al., 2019 Balezeau et al., 2020 Rocchi et al., 2021	/

5 Table 1.: Summary of the main work analysed in the present review, in regard to the contrasting  
6 descriptions of the Arcuate Fasciculus (AF) in tract-tracer and diffusion MRI studies. Six features  
7 are highlighted as reported or not reported: Broca's terminations in monkeys, temporal  
8 terminations into the monkeys' auditory cortex in the temporal lobe, expanded terminations into



the human's middle and anterior temporal lobe (see e.g., Eichert et al., 2019), dominance of the Ventral Pathway over the Dorsal Pathway in monkeys, lateralisation in chimpanzee brains and lateralisation in monkey brains. “/”: No data available.

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