



**Alain Prochiantz**

Professeur

**Evolution du cerveau :  
les singularités de l'animal  
humain**

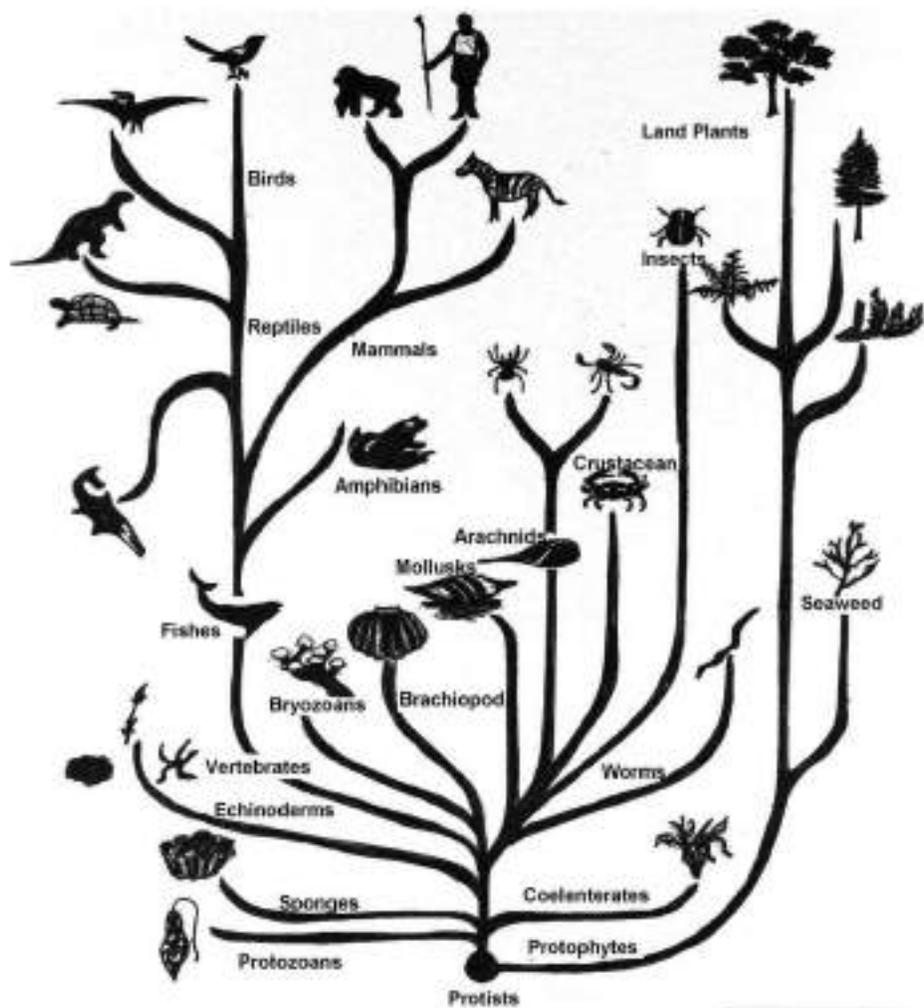
**Journées LIT Expert**

*Jeudi 13 octobre 2022*



# Evolution du cerveau: les singularités de l'animal humain

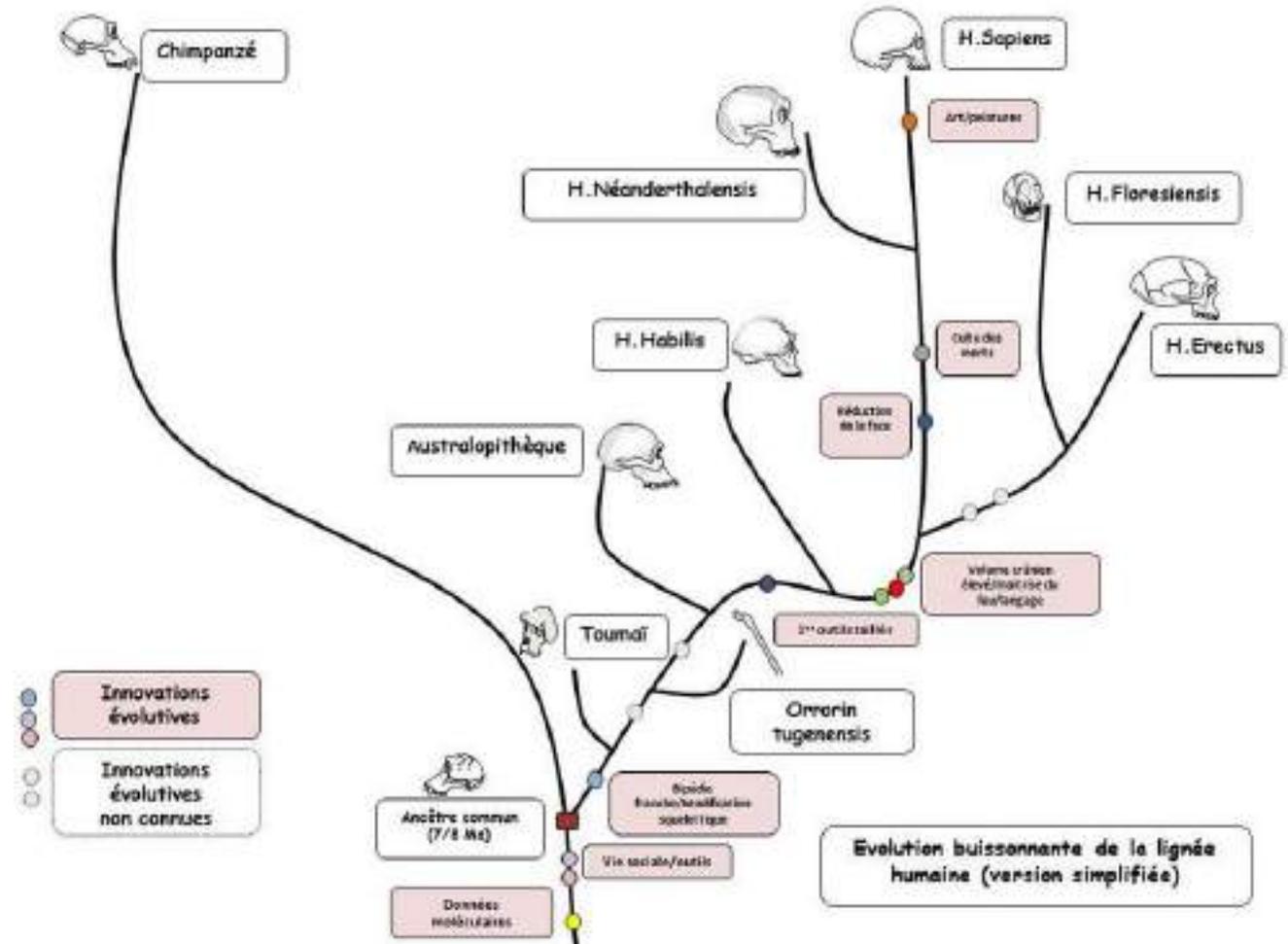
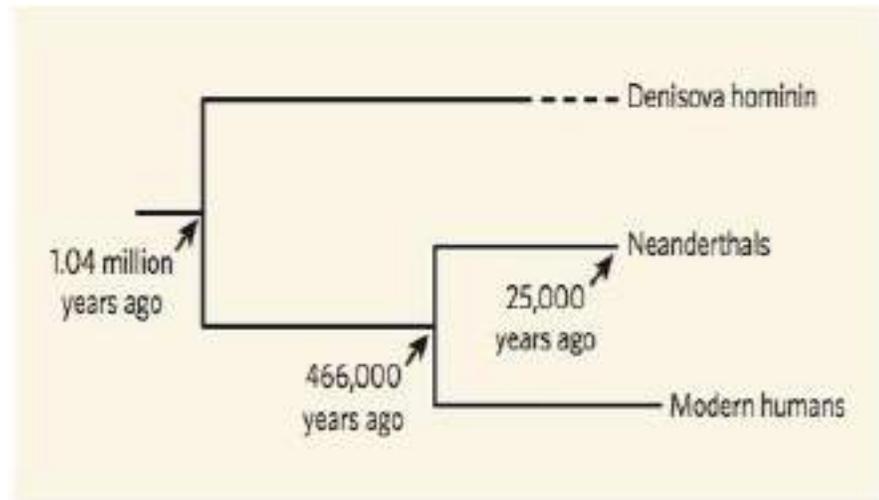
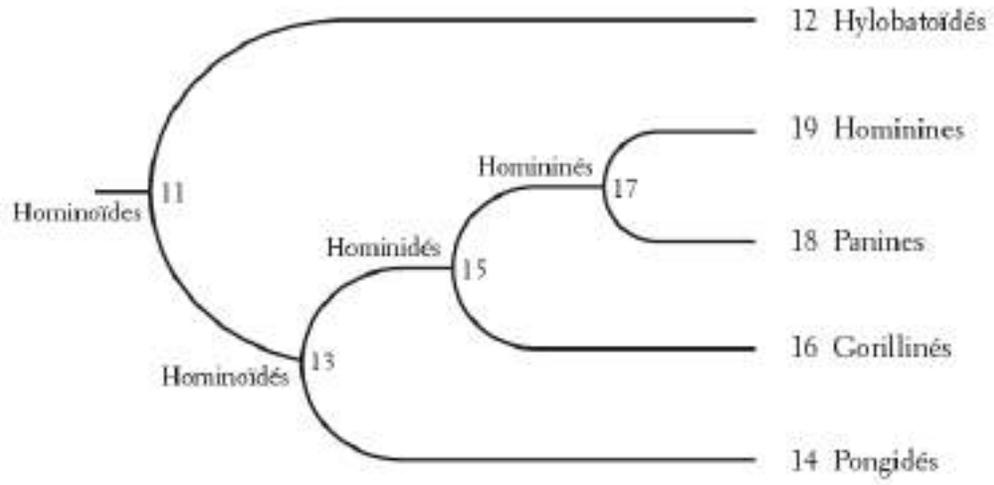


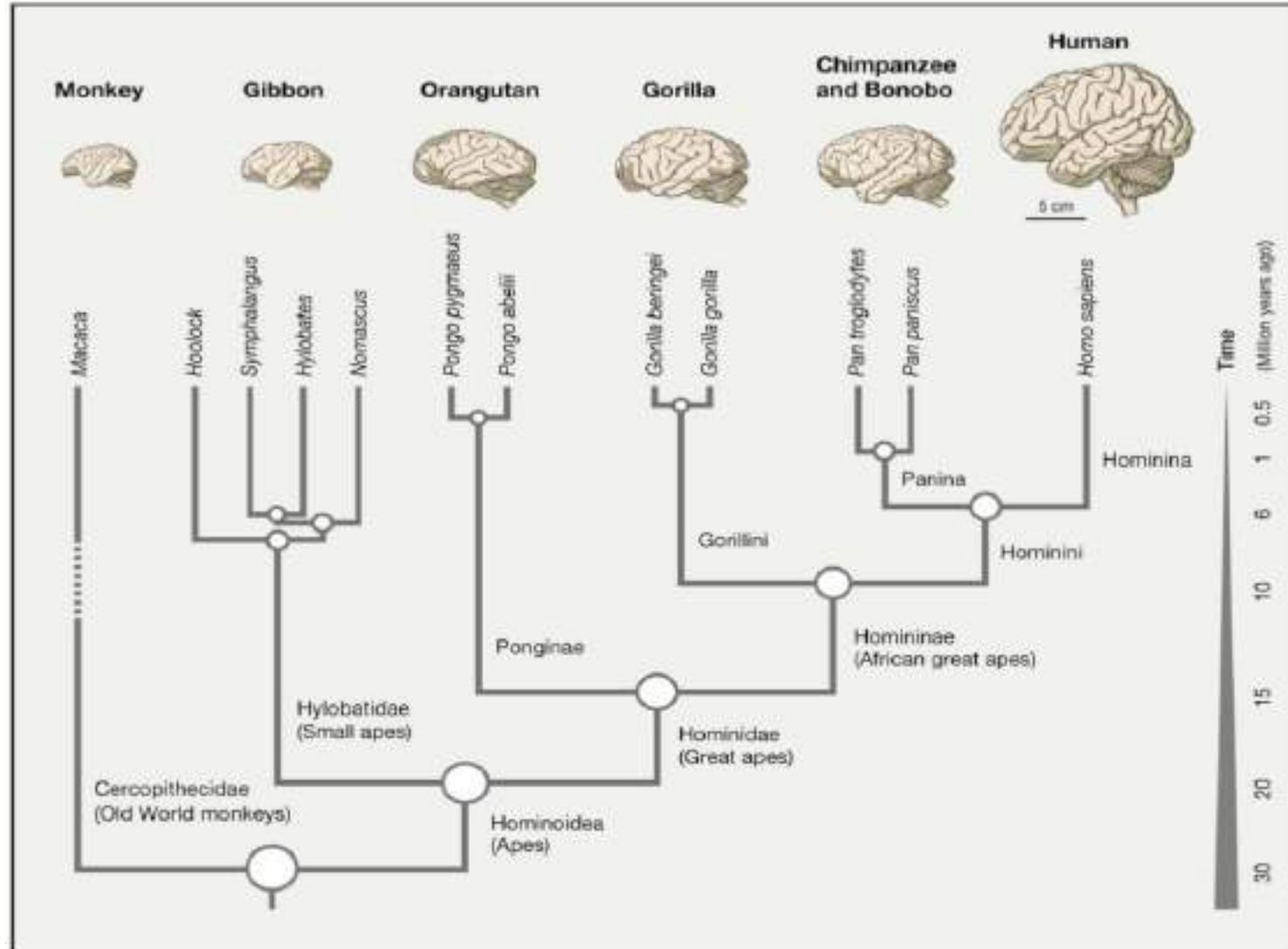


Ère	Époque	Âge (années)	Événements
Quaternaire	Holocène		Agriculture
	Pléistocène	126 000 2,6 millions	Évolution des humains modernes
Tertiaire	Pliocène	3,6 millions	Lucy ←
		5,3 millions	
	Miocène	7 millions	Séparations des lignées humaines et chimpanzés/bonobos
		23 millions	
	Oligocène	28 millions	Isolement du continent antarctique
		34 millions	
Éocène	38 millions	Nombreuses nouvelles espèces de petits mammifères	
	56 millions		
Paléocène	59 millions	Premiers primates	
	66 millions		

« Donc nous avons attribué à l'homme un pedigree d'une longueur prodigieuse, mais pas, pourrait-on dire, de noble qualité. Le monde, cela a souvent été dit, semble s'être préparé de longue main à la venue de l'homme ; et cela, en un sens est strictement vrai, car il doit sa naissance à une longue lignée de progéniteurs. Si un seul des liens de cette chaîne n'avait jamais existé, l'homme n'aurait pas été exactement ce qu'il est aujourd'hui. A moins que nous ne nous aveuglions volontairement, nous pouvons, avec nos connaissances actuelles, reconnaître approximativement notre parenté ; et nous n'avons nulle raison d'en concevoir de la honte. L'organisme le plus humble est une chose très supérieure à la poussière inorganique sous nos pieds ; et personne avec un esprit non biaisé ne peut étudier une créature vivante, quelque humble qu'elle soit, sans être frappé d'enthousiasme devant le merveilleux de sa structure et de ses propriétés. »

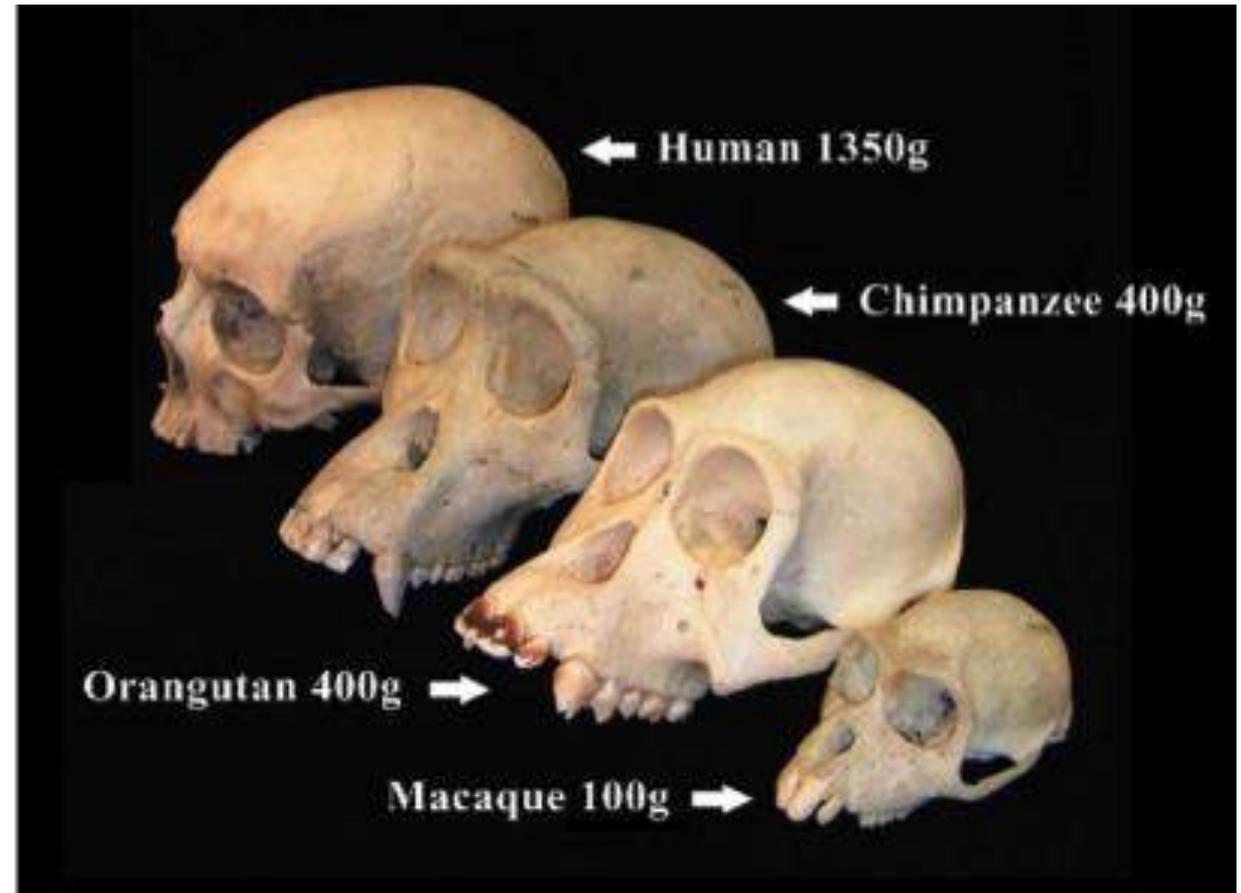
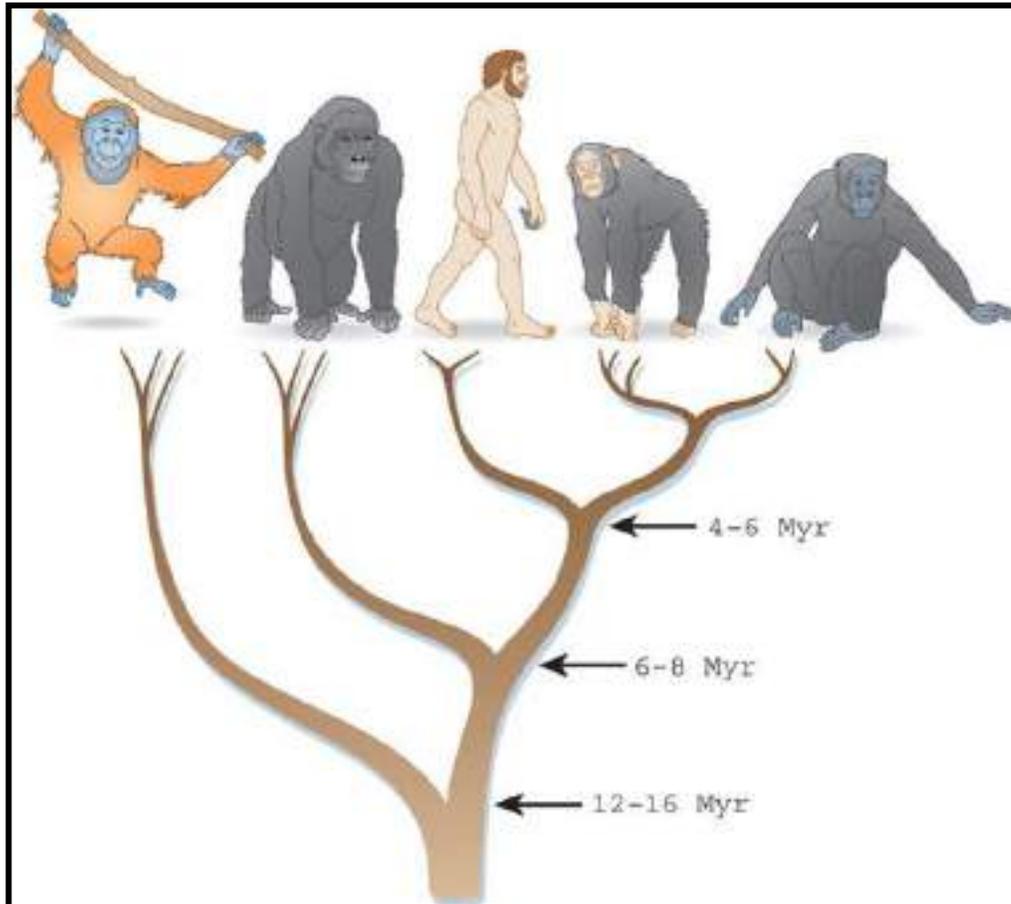
Darwin, The descent of Man

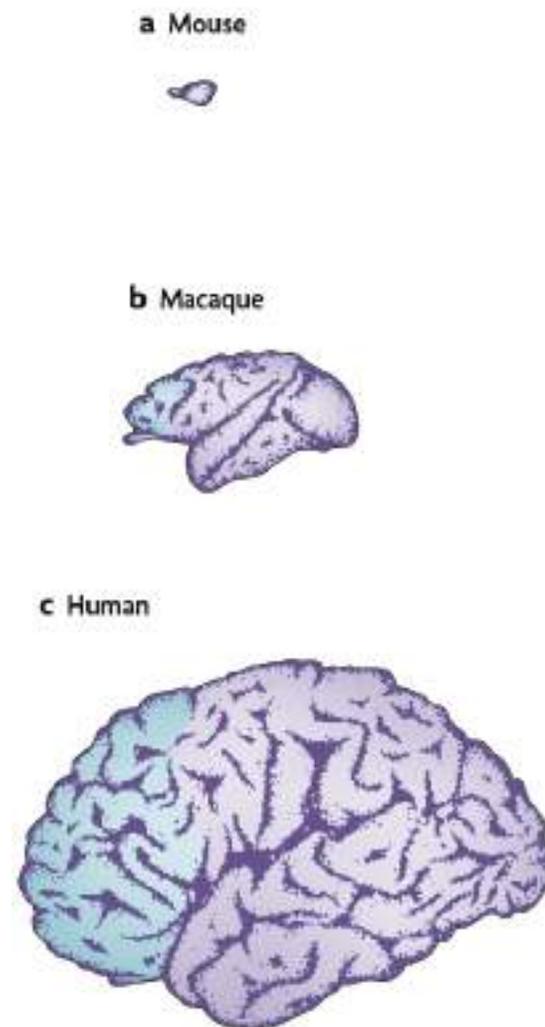
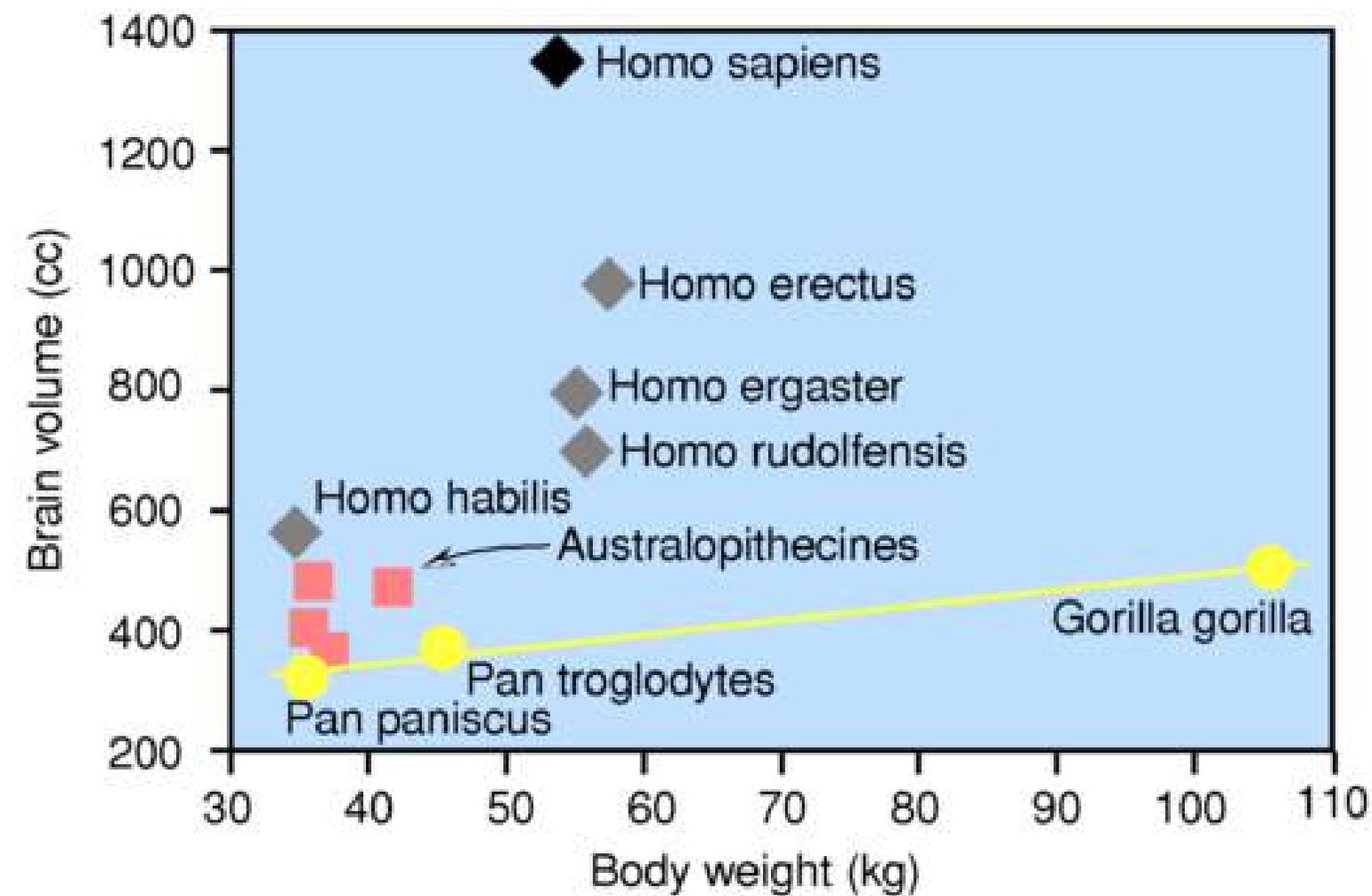




Evolution of the Human Nervous System  
 Function, Structure, and Development.

André M M Sousa et al.  
 Cell, 2017 vol. 170 (2) pp. 226-247



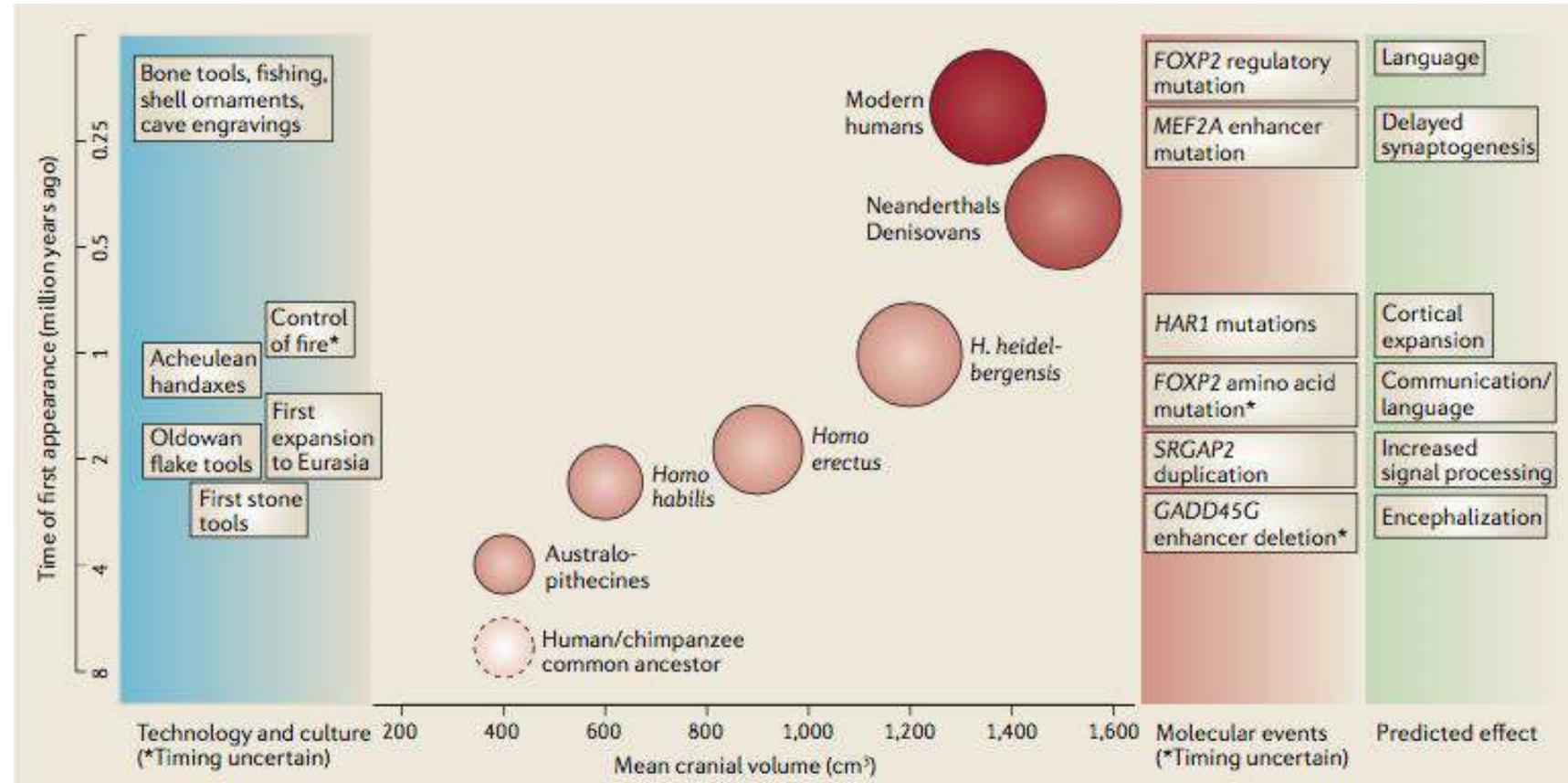


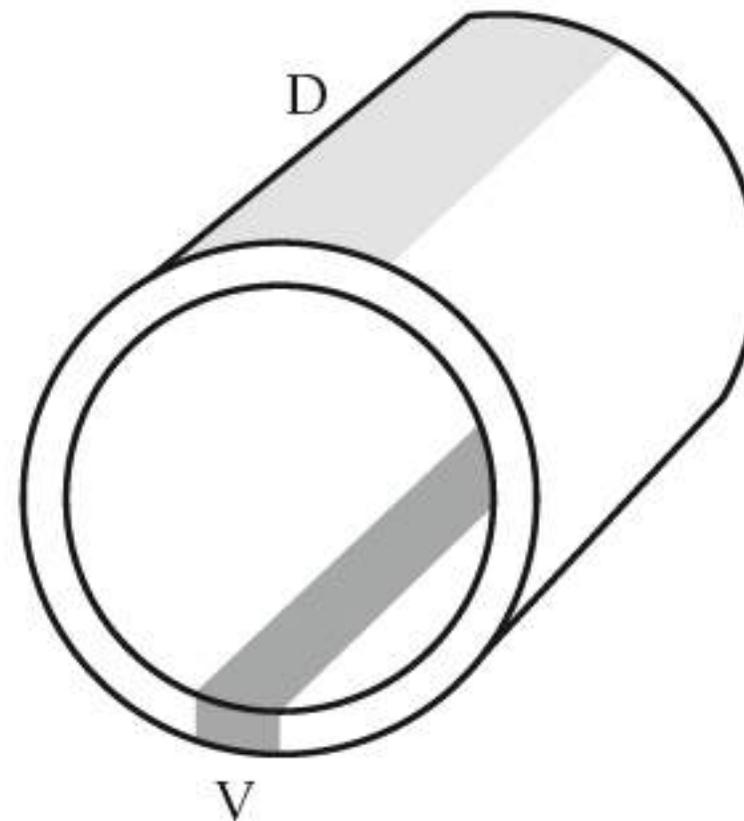
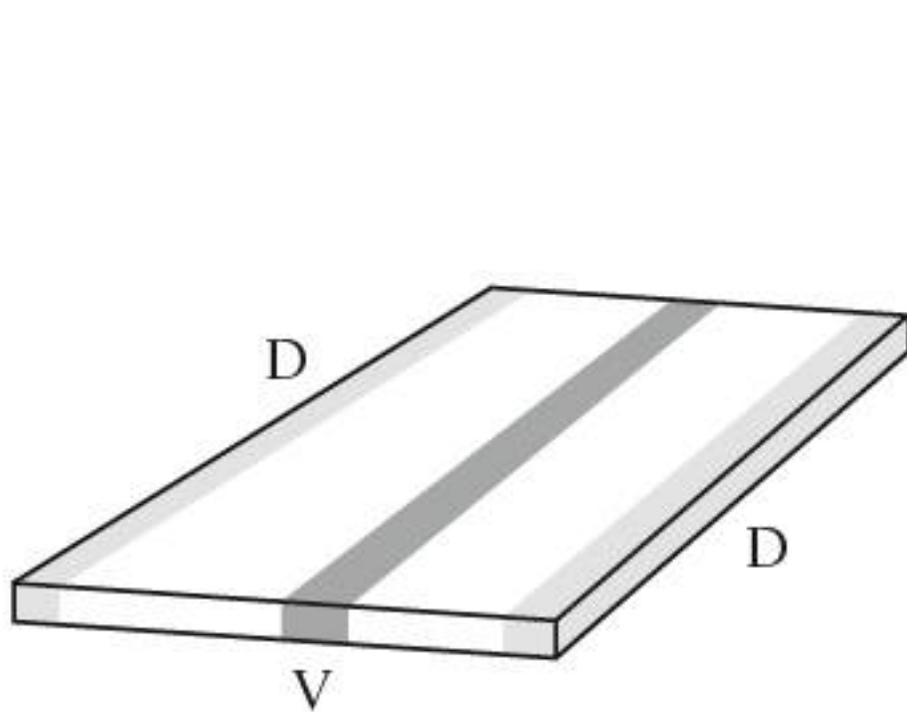
# Human brain evolution: transcripts, metabolites and their regulators

Mehmet Somel<sup>1,2</sup>, Xiling Liu<sup>1</sup> and Philipp Khaitovich<sup>1,5</sup>

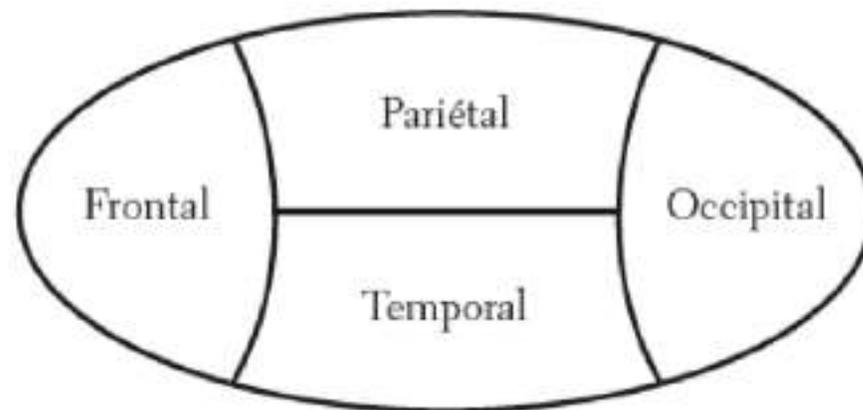
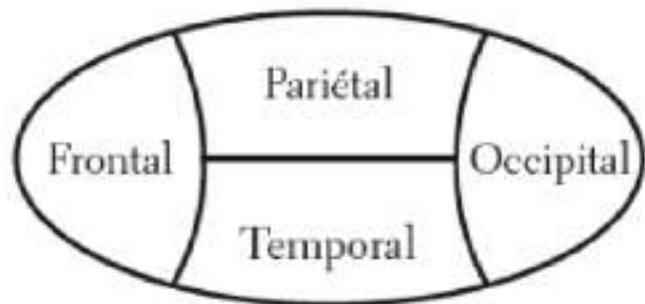
112 | FEBRUARY 2013 | VOLUME 14

[www.nature.com/reviews/neuro](http://www.nature.com/reviews/neuro)

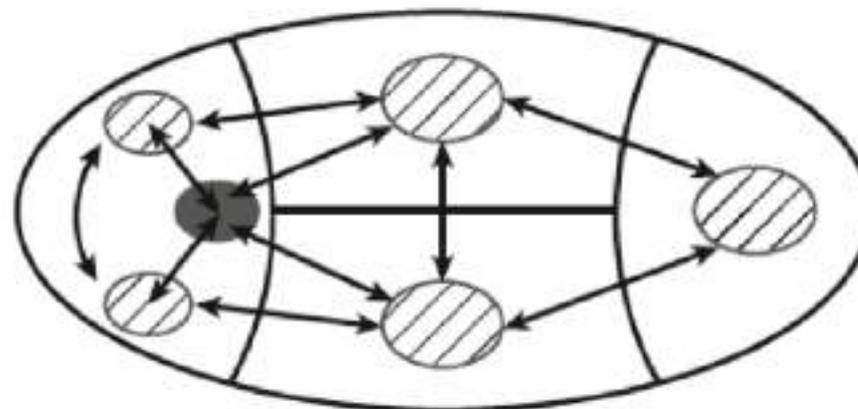
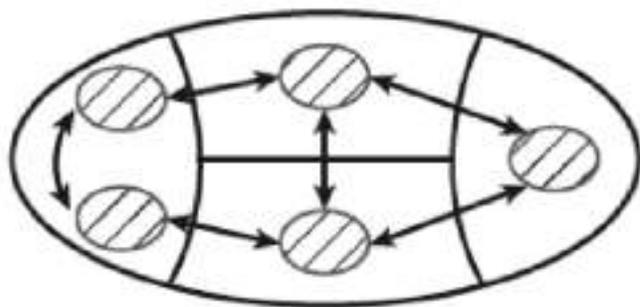




A.



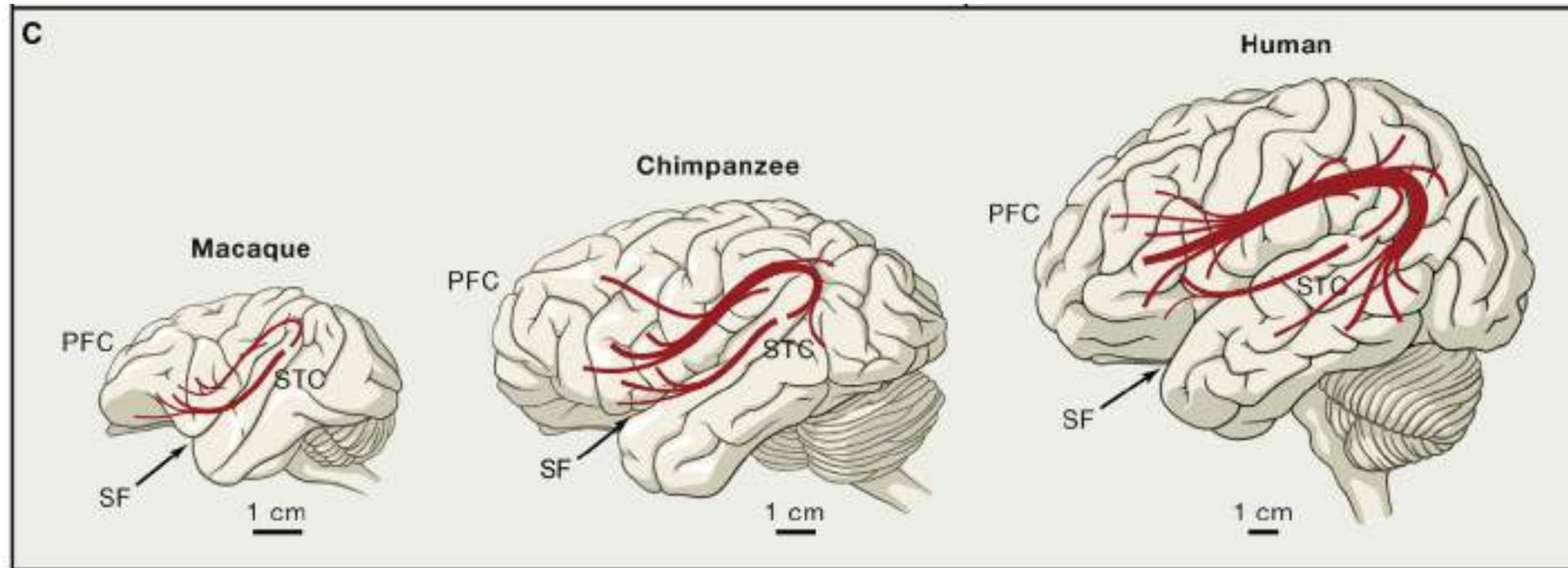
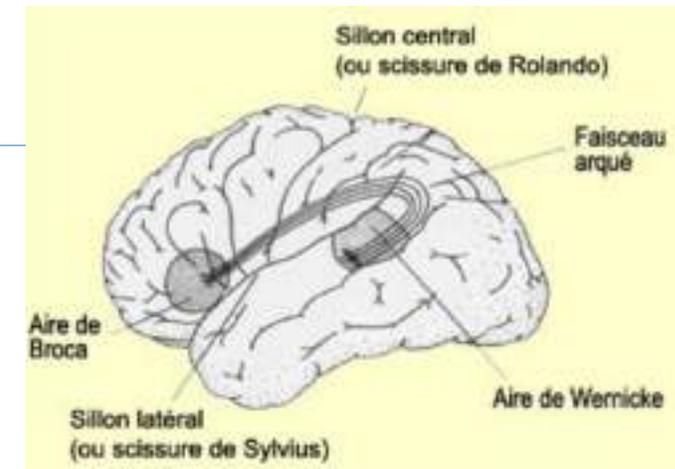
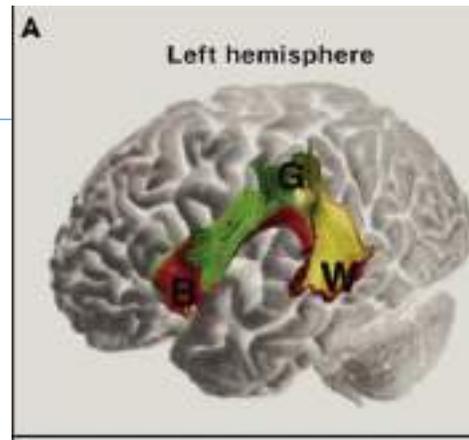
B.



# Evolution of the Human Nervous System Function, Structure, and Development.

André M M Sousa et al.

Cell, 2017 vol. 170 (2) pp. 226-247



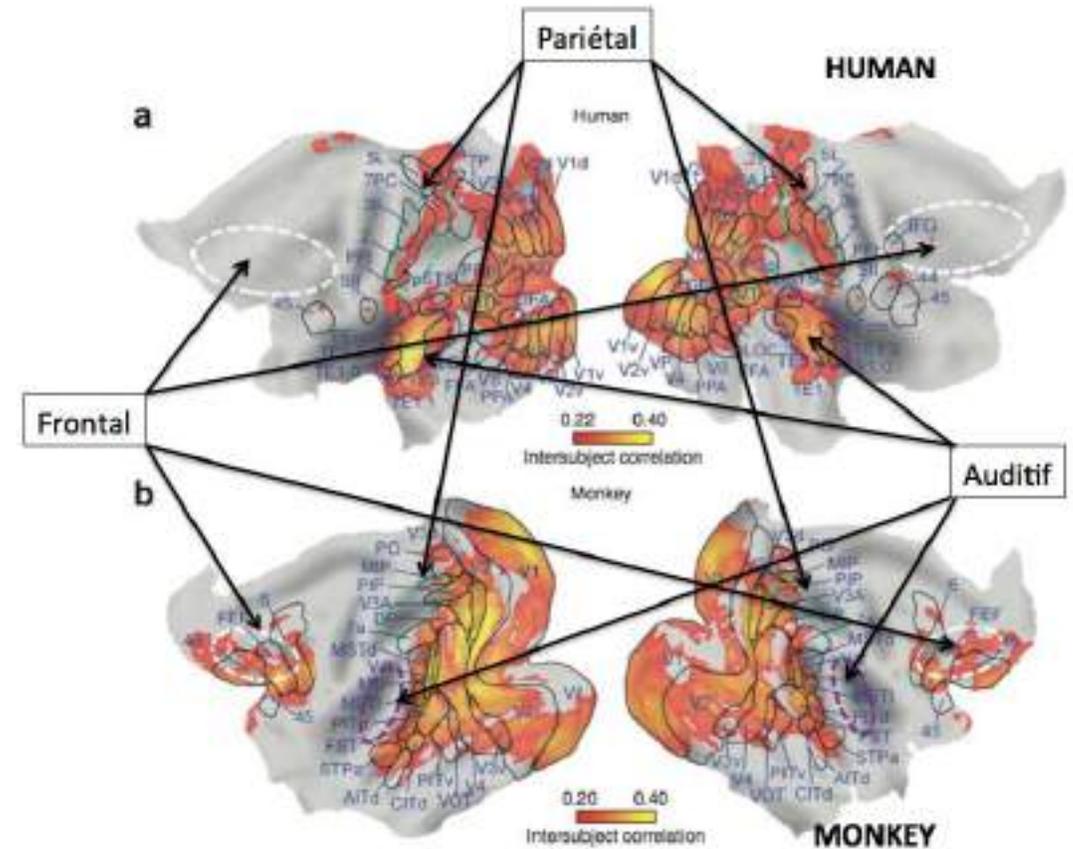
**Figure 3. Language-Related Pathways Are Strongly Lateralized and Modified in Humans**

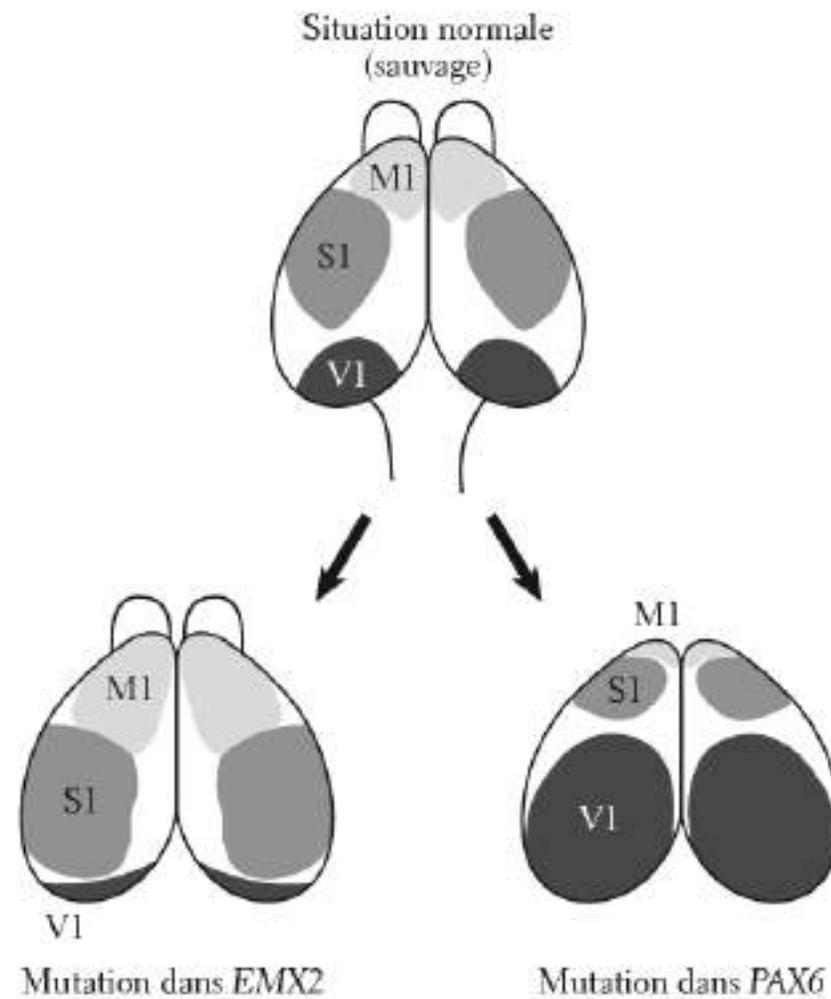
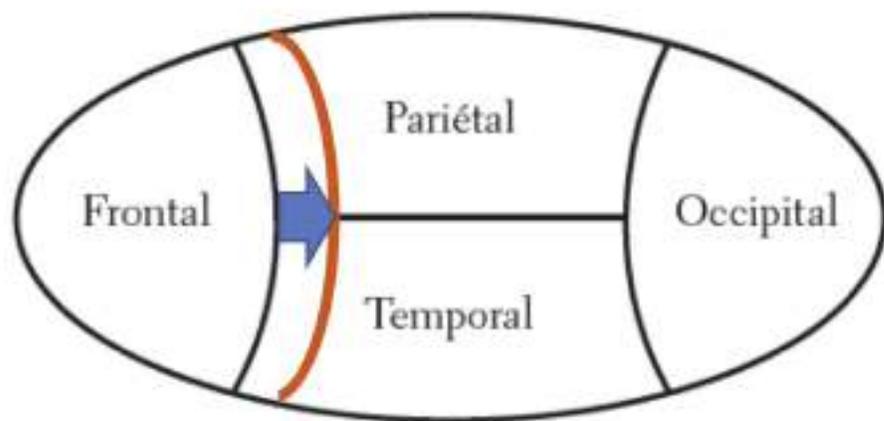
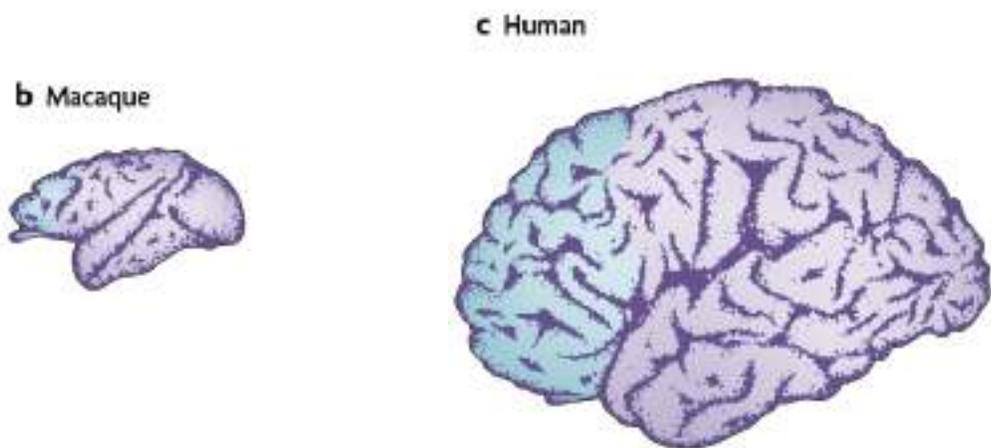
# Establishing homology between monkey and human brains

Tor D Wager & Tal Yarkoni



**Figure 1** | The ISAC method. Humans and monkeys watch the same film as their brains are scanned with fMRI. The time course of fMRI activity during viewing is extracted from 'seed' regions in human participants and correlated with the fMRI signal in monkeys, and vice versa, with an adjustment made for interspecies differences in vascular hemodynamics. After statistical thresholding, the resulting maps in each species show areas that are potentially homologous with the targeted seed region in the other species.



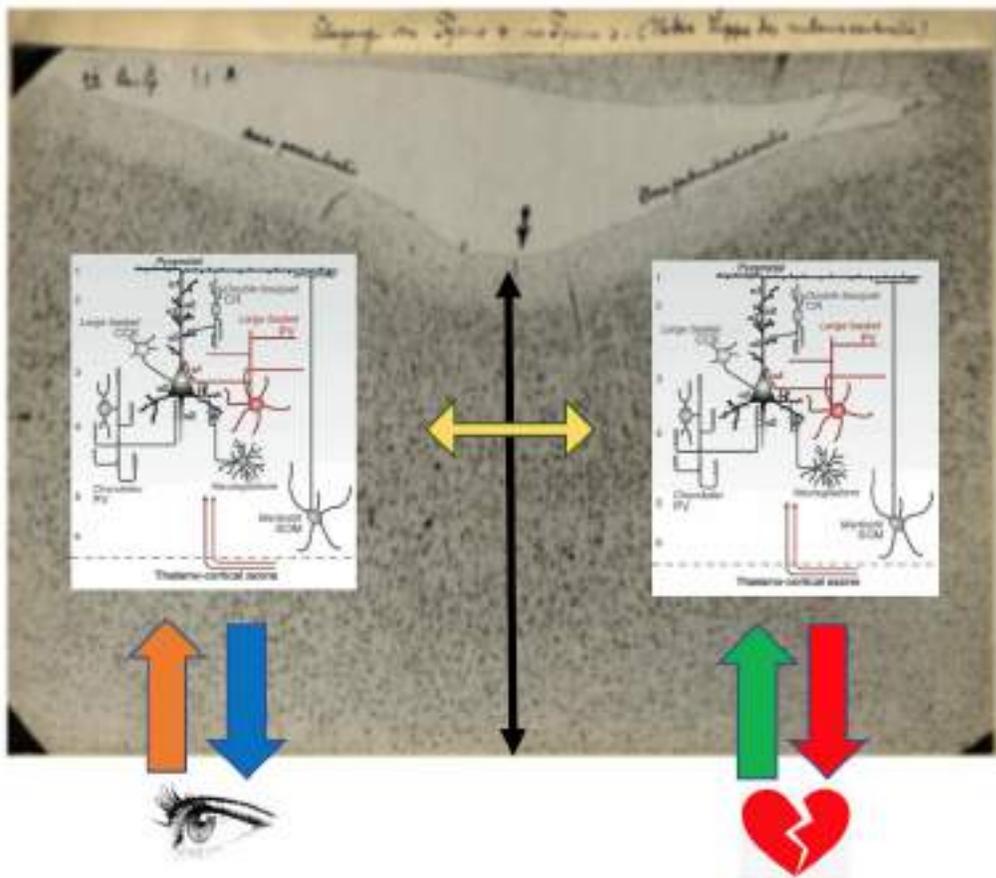


# Centenary of Brodmann's map — conception and fate

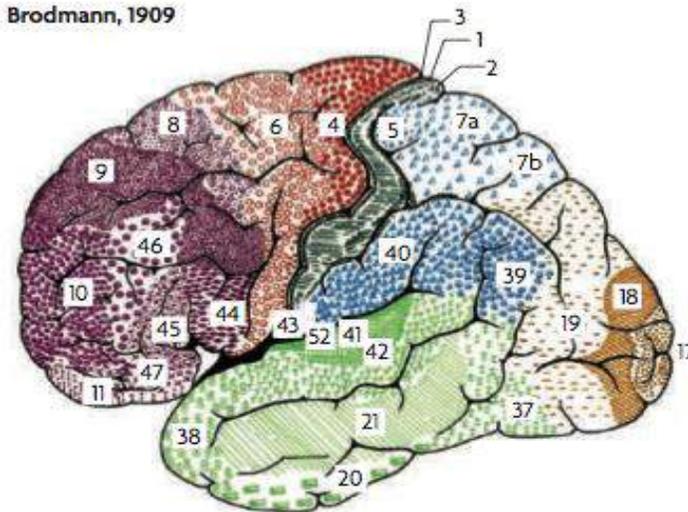
Karl Zilles and Katrin Amunts

NATURE REVIEWS | NEUROSCIENCE

VOLUME 11 | FEBRUARY 2010 | 219



Brodmann, 1909

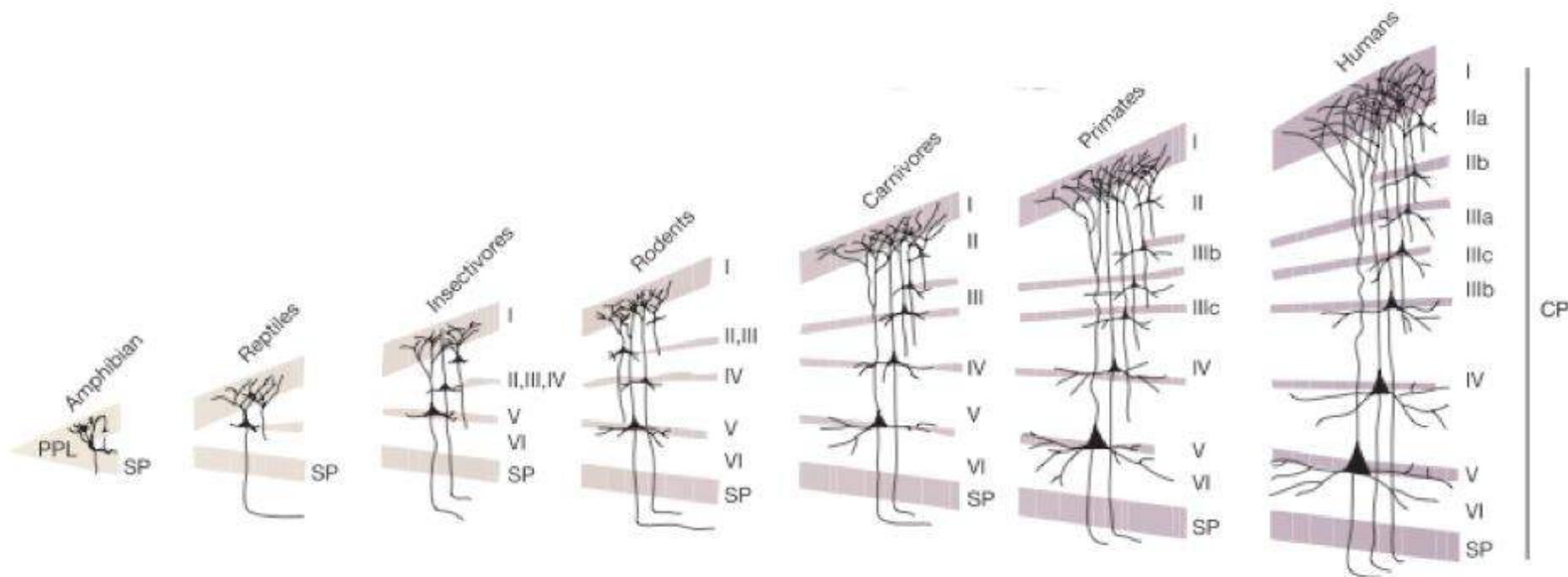


Augmentation de la taille  
 Augmentation du nombre des aires  
 Déplacement des bord  
 Changement des circuits

# PROGRESS

## Molecular insights into human brain evolution

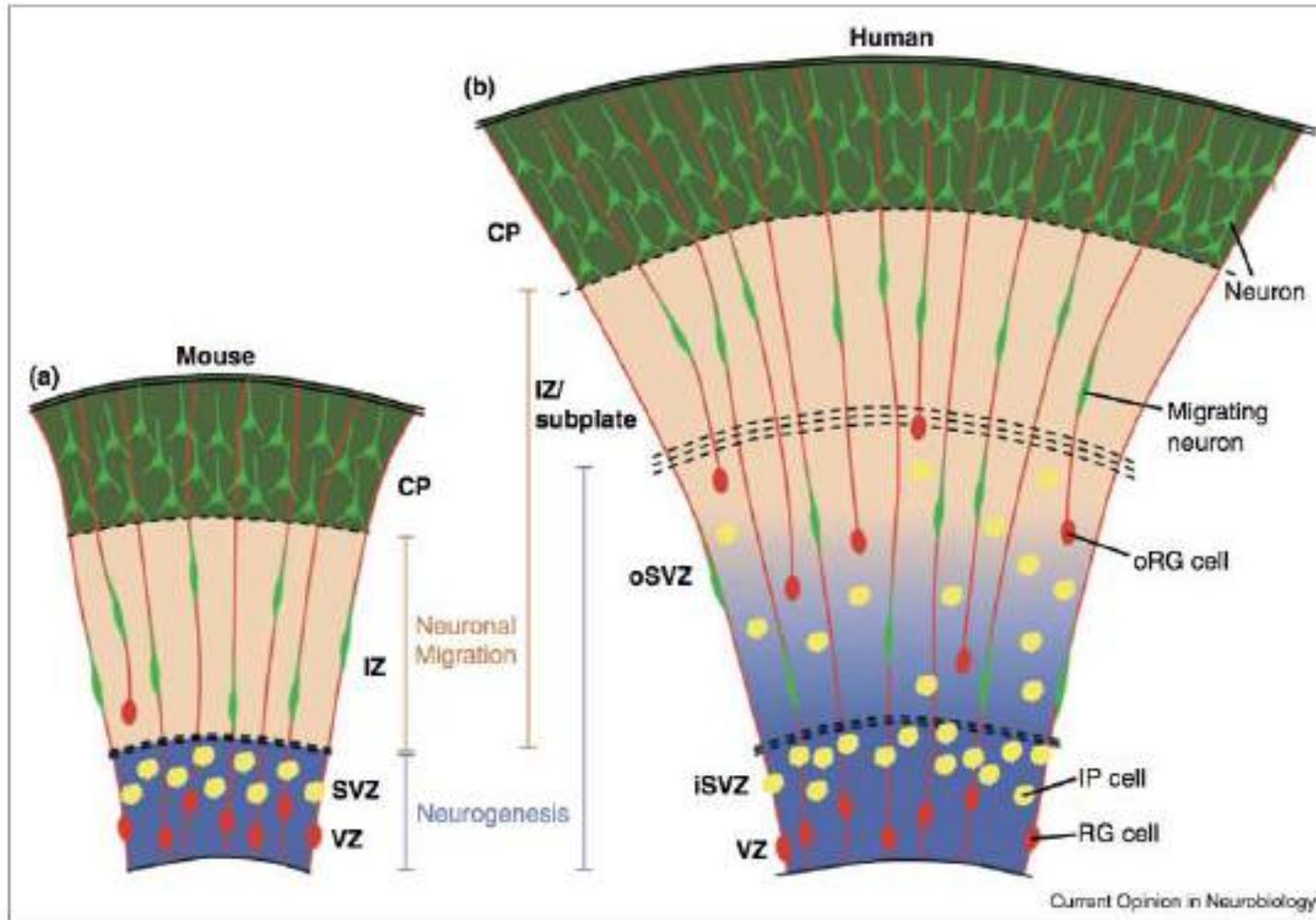
Robert Sean Hill<sup>1</sup> & Christopher A. Walsh<sup>1</sup>



# OSVZ progenitors in the human cortex: an updated perspective on neurodevelopmental disease

Bridget E LaMonica<sup>1,2,3</sup>, Jan H Lui<sup>1,2</sup>, Xiaoqun Wang<sup>1,2</sup> and Arnold R Kriegstein<sup>1,2</sup>

Current Opinion in Neurobiology 2012, 22:747–753



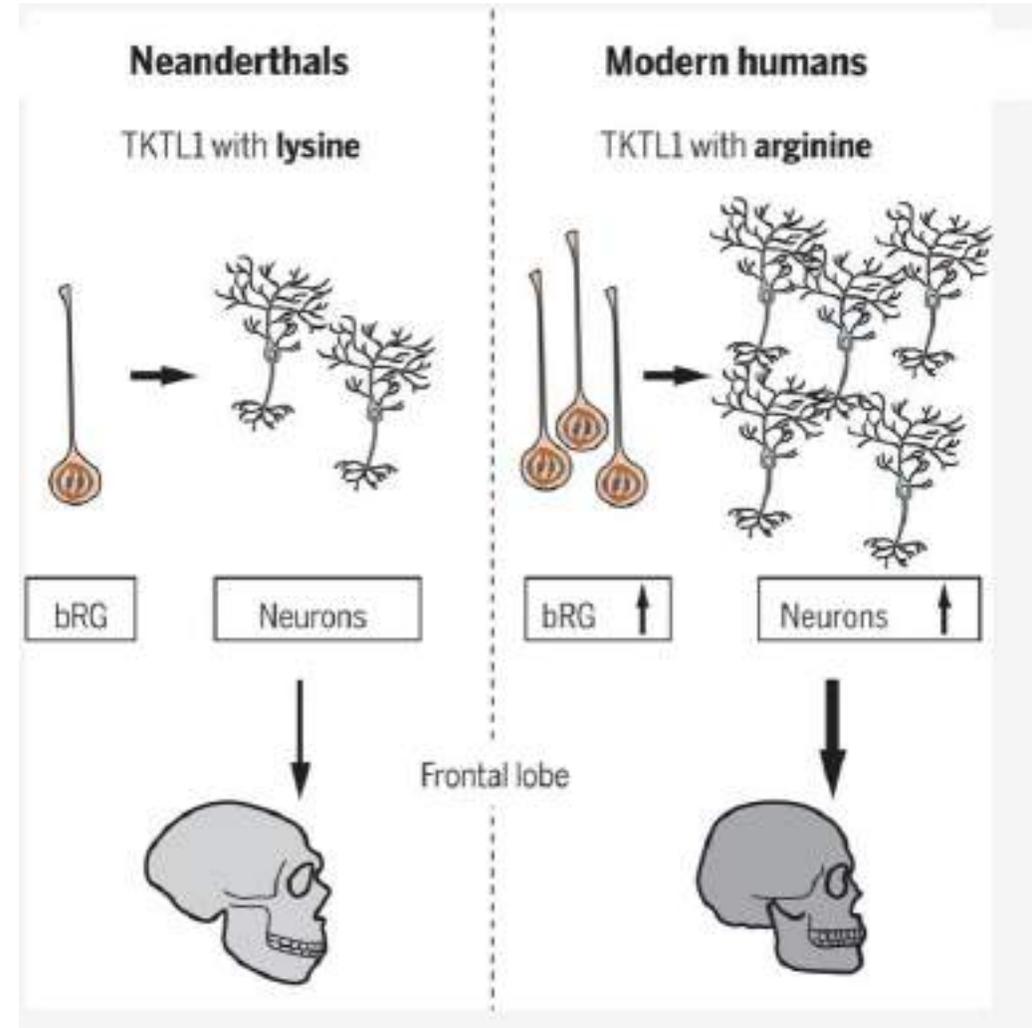
# Science

SCIENCE > VOL. 377, NO. 6611 >

## Human TKTL1 implies greater neurogenesis in frontal neocortex of modern humans than Neanderthals

ANNELINE HINGSON, LEI XING, TAKASHI NAMBA, RENE KALERIC, JILA PEYER, CHRISTINA FUGSTER CEDENA, SOFIA TRANKWY, KATRIN REPE, STRIMMER REIMERS, TOMELAW MARICIC, RACHMADEBIBACI, ISOLINE HUMBERGER, SWANTE PASARO, AND WELANGLD. HUTTEN

[Authors info & Affiliations](#) fewer

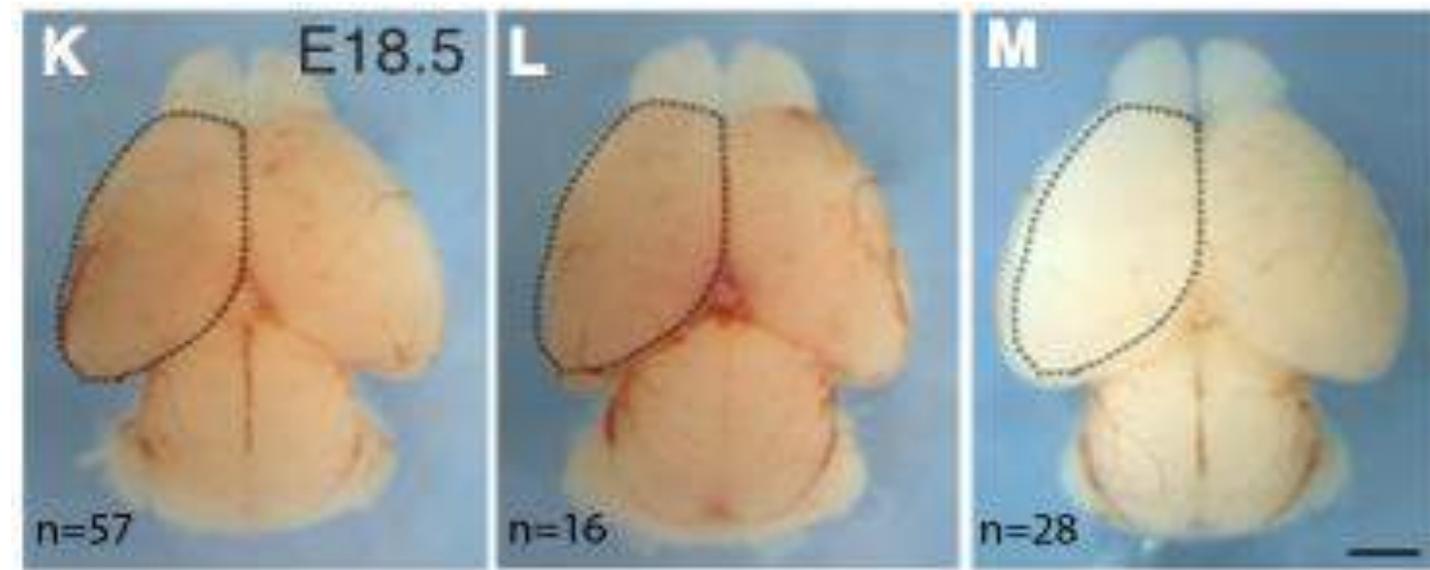
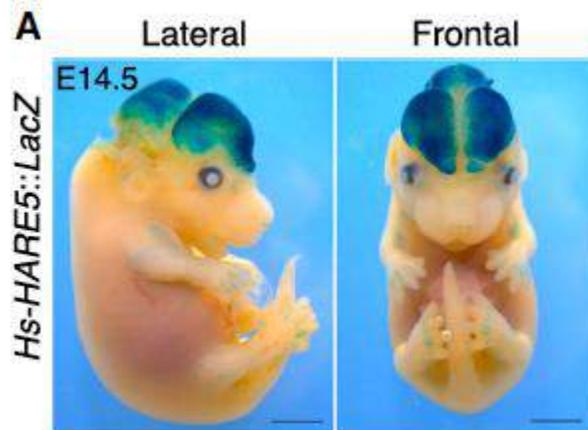
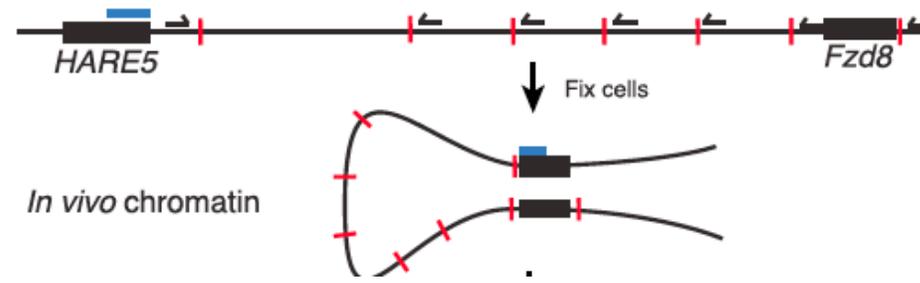


## HAREs ou Régions accélées (en nombre de mutations) chez l'humain

Human-chimpanzee differences in a FZD8 enhancer alter cell-cycle dynamics in the developing neocortex.

J Lomax Boyd et al.

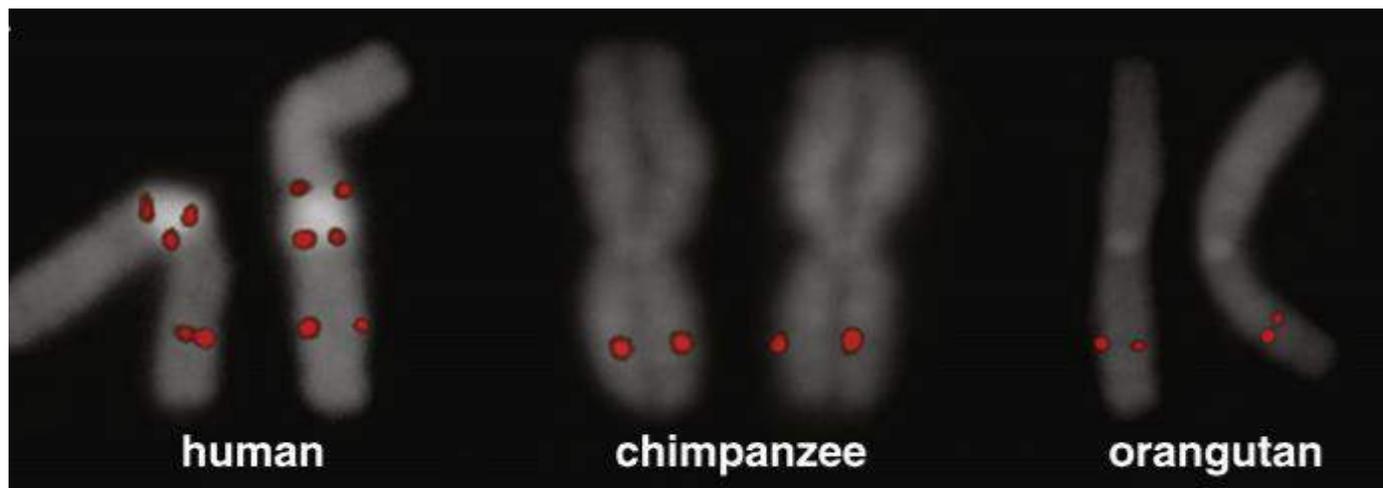
Curr Biol, 2015 vol. 25 (6) pp. 772-779



Evolution of human-specific neural SRGAP2 genes by incomplete segmental duplication.

Megan Y Dennis et al.  
Cell, 2012 vol. 149 (4) pp. 912-922

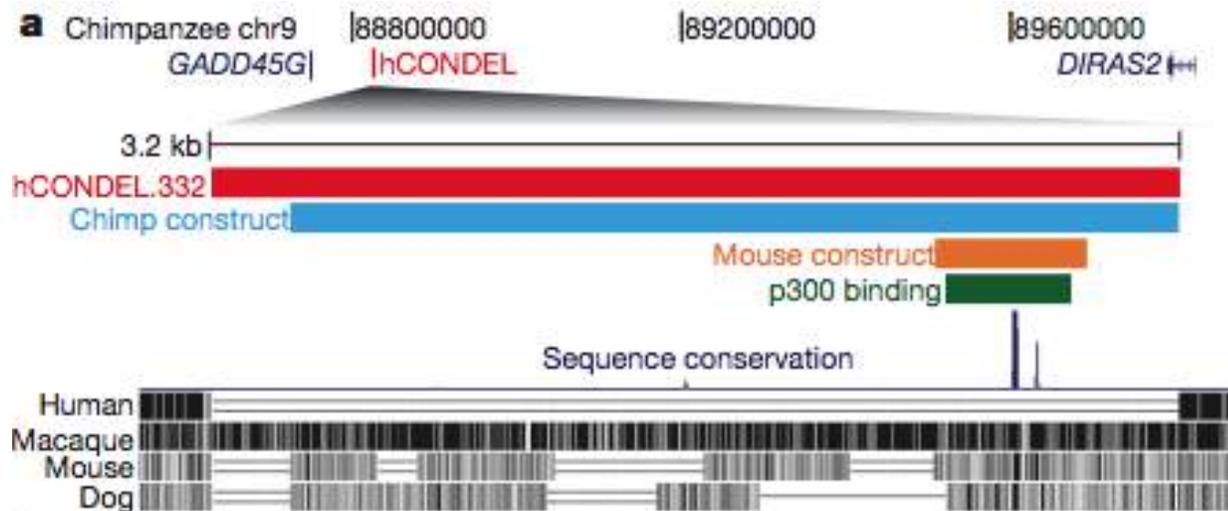
Duplication



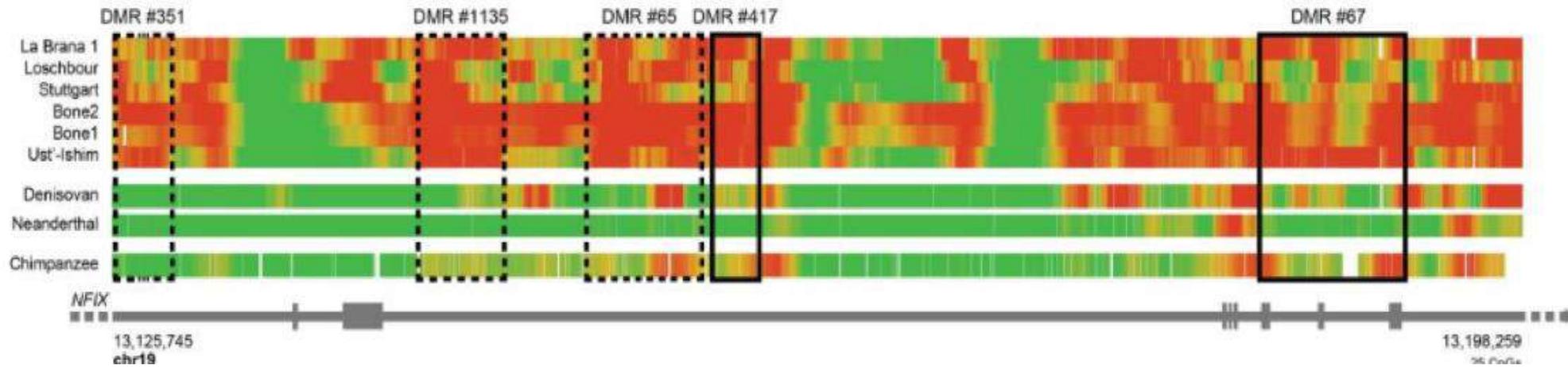
Mclean CY, Reno PL, Pollen AA, Bassan AI, Capellini TD, Guenther C, Indjeian VB, Lim X, Menke DB, Schaar BT, Wenger AM, Bejerano G, Kingsley DM

Nature  
2011 vol. 471 (7337) pp. 216-9

Délétion.

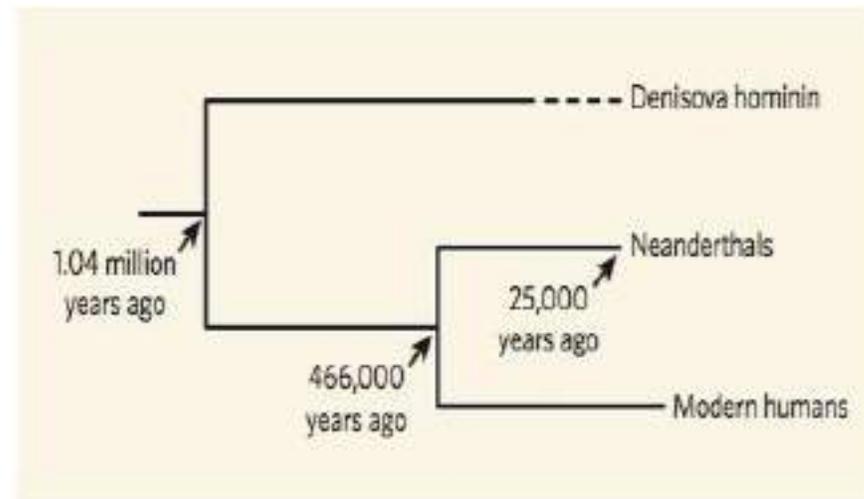


Human-specific loss of regulatory DNA and the evolution of human-specific traits



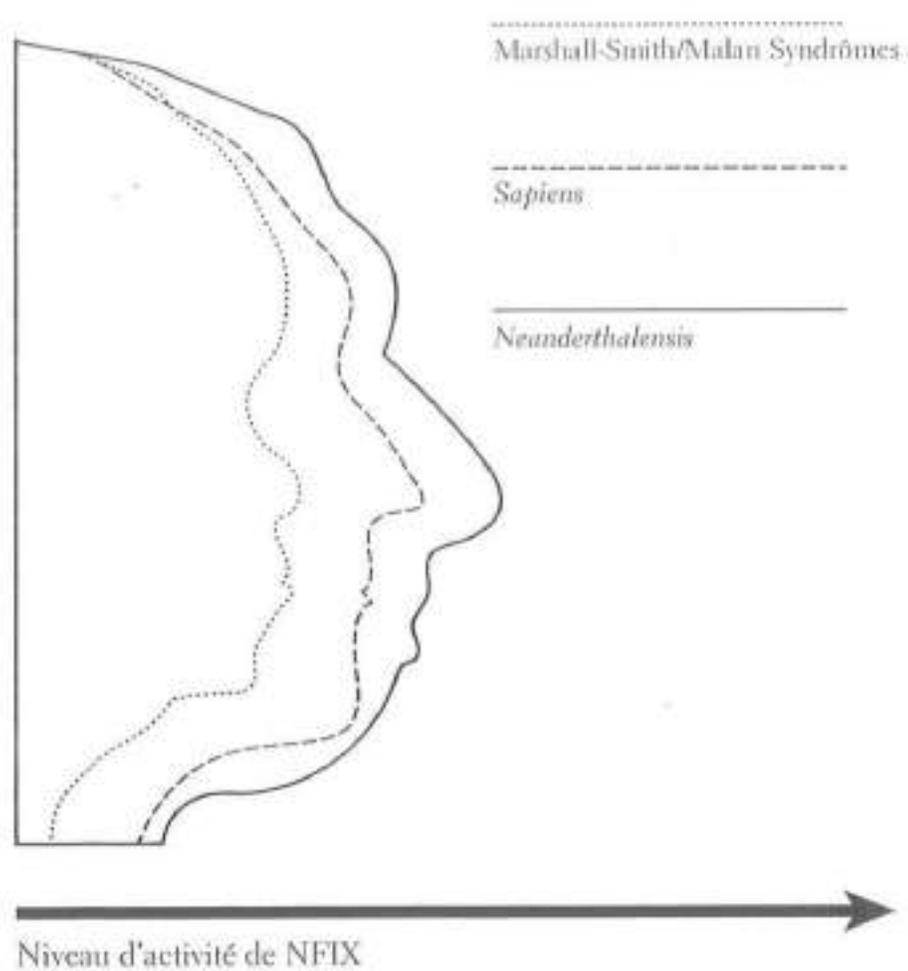
**Extensive Regulatory Changes in Genes Affecting Vocal and Facial Anatomy Separate Modern from Archaic Humans**

David Gokhman, Lily Agranon-Tamir, Genevieve Houtman, Raquel Garcia-Perez, Malika Nizam-Ibrahima, Swapan Mallick, Maria Nieves-Colón, Heng Li, Songxi Alpaalen-Roodenberg, Mario Novak, Hongchang Gu, Manuel Ferrando-Bernal, Pere Galabert, Iddi Lipande, Ivanela Kondova, Ronald Berthrop, Eilon E. Quillen, Alexander Heitsch, Anne C. Stone, Anne E. Pusey, Deus Mjunga, Leonid Kandel, Meir Liebergall, Moris E. Prada, Julio M. Yáñez, Kay Prüfer, Johannes Krause, Benjamin Yalcin, Svante Pääbo, Raz Pihasi, Carlos Lalueza-Fox, David Reich, Tomas Marques-Bonet, Eran Meshorer, Liran Carmel  
doi: <https://doi.org/10.1101/106955>



**Table 1.** DMGs that affect the voice and the larynx.

DMG	Associated phenotype	Chr
ALPL	Abnormality of the voice	1
AHDC1	Laryngomalacia	1
AHDC1	Laryngomalacia	1
SATB2	Abnormality of the voice	2
SPEG	Dysphonia	2
COLQ	Weak cry	3
TGFBR2	Abnormality of the voice	3
TGFBR2	Abnormality of the voice	3
TGFBR2	Abnormality of the voice	3
POC1A	High pitched voice	3
PLXND1	Abnormality of the voice	3
SH3BP2	Abnormality of the voice	4
BDHA	Hoarse voice, loss of voice, vocal cord paralysis	5
GLI3	Laryngeal cleft	7
CHD7	Abnormality of the voice, Laryngomalacia	8
HNRNPA1	Bowing of the vocal cords, hoarse voice	12
TRPV4	Vocal cord paresis	12
MEIS2	Laryngomalacia	15
ACAN	Hoarse voice	15
CREBBP	Laryngomalacia	16
CREBBP	Laryngomalacia	16
XYLT1	High-pitched voice	16
WWOX	Abnormality of the voice	16
WWOX	Abnormality of the voice	16
SOX9	Laryngomalacia	17
SOX9	Laryngomalacia	17
GNAL	Laryngeal dystonia	18
NFIX	Laryngomalacia	19
NFIX	Laryngomalacia	19
PCLO1	High-pitched voice	19
RIN2	High-pitched voice	20
TBX1	Abnormality of the voice, nasal speech	22



## Larynx et cordes vocales †

langue †

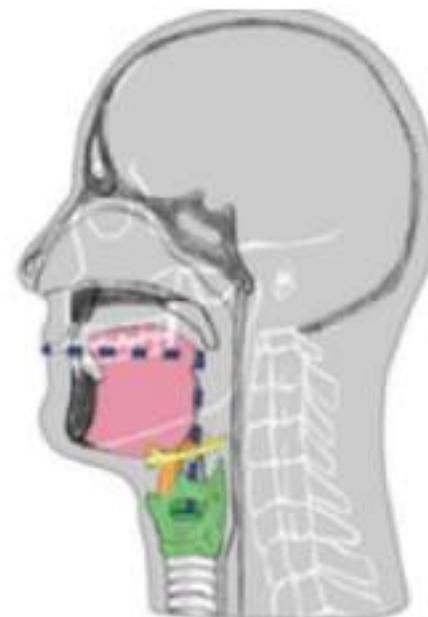
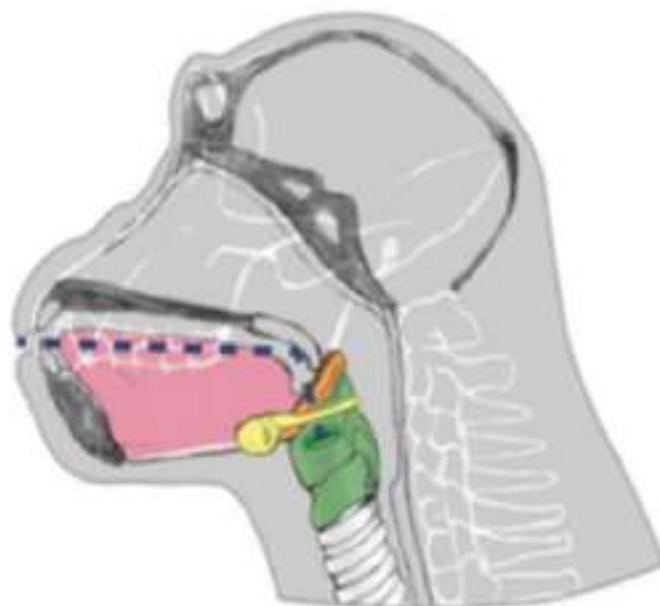
Os hyoïde †

épigllotte †

David Gokhman<sup>1</sup>, Lily Agronai-Tamir<sup>1,2</sup>, Genevieve Houman<sup>1,4</sup>, Rapel Garcia-Pérez<sup>5</sup>, Malka Nissim-Rafinia<sup>1</sup>, Swapan Mallick<sup>6,7,8</sup>, María A. Nieves-Collín<sup>1,4</sup>, Hongchang Gu<sup>9</sup>, Manuel Ferrando-Bernal<sup>1</sup>, Pere Gelabert<sup>1</sup>, Iddi Lipende<sup>9</sup>, Ivanela Kondova<sup>10</sup>, Ronald Bontrop<sup>10</sup>, Ellen E. Quillen<sup>11</sup>, Alexander Meissner<sup>6,12,13</sup>, Anne C. Stone<sup>14,15</sup>, Anne E. Pusey<sup>15</sup>, Deus Mjunga<sup>9</sup>, Leonid Kandel<sup>16</sup>, Meir Liebergall<sup>10</sup>, María E. Prada<sup>17</sup>, Julio M. Vidal<sup>18</sup>, Kay Prüfer<sup>19</sup>, Johannes Krause<sup>20</sup>, Benjamin Yakir<sup>2</sup>, Svanne Pääbo<sup>19</sup>, David Reich<sup>6,7,8</sup>, Carles Lahozza-Fox<sup>2</sup>, Tomas Marques-Bonet<sup>6,12,21</sup>, Eran Meshorer<sup>1,22\*</sup>, Liran Carmel<sup>1,7</sup>

†

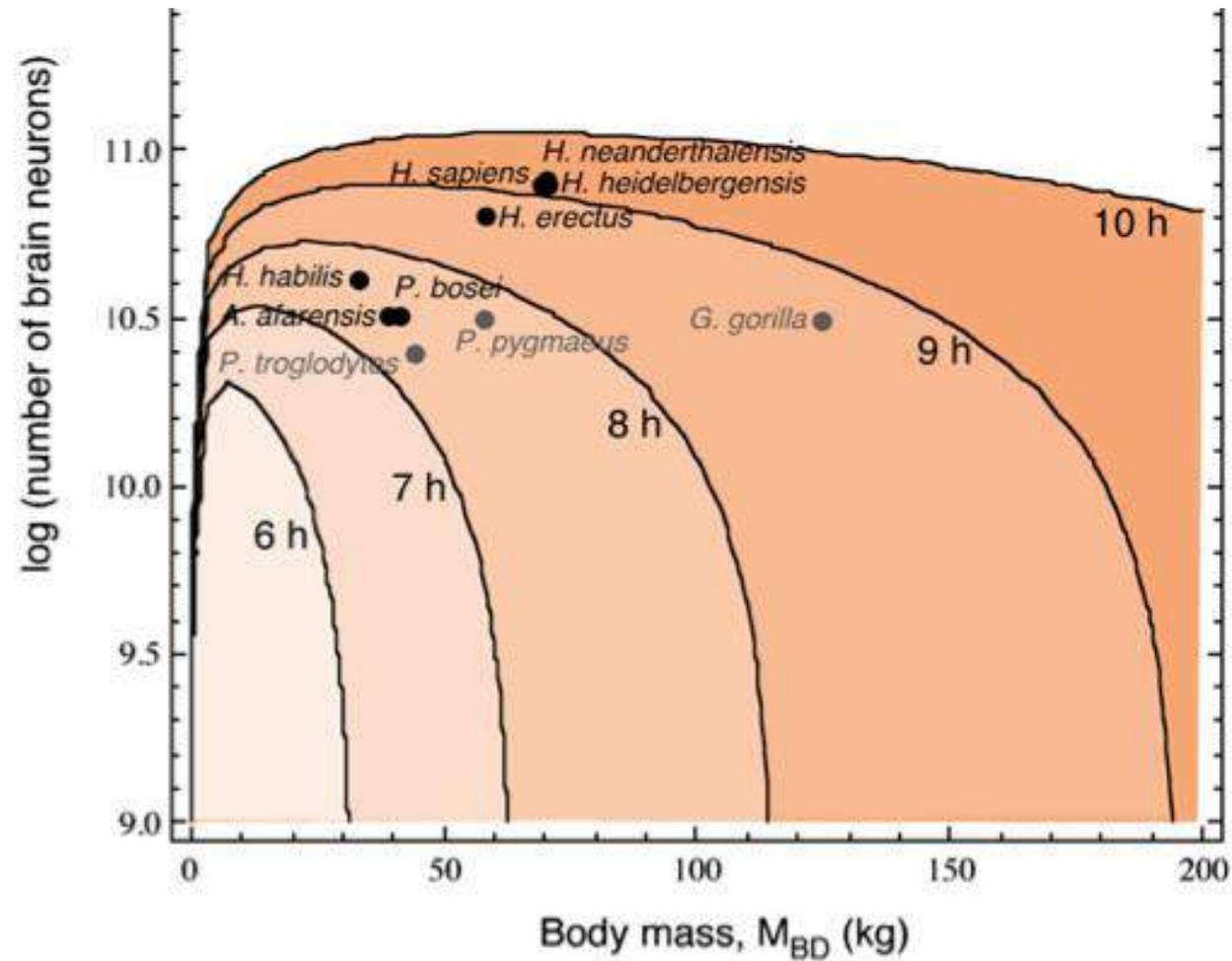
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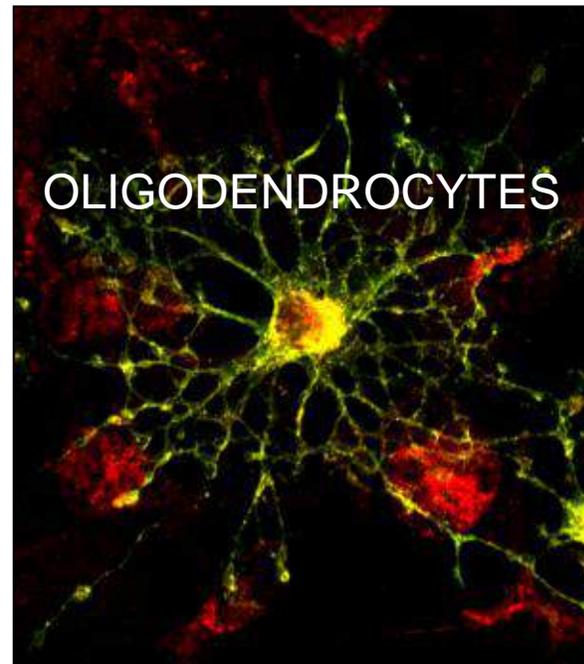
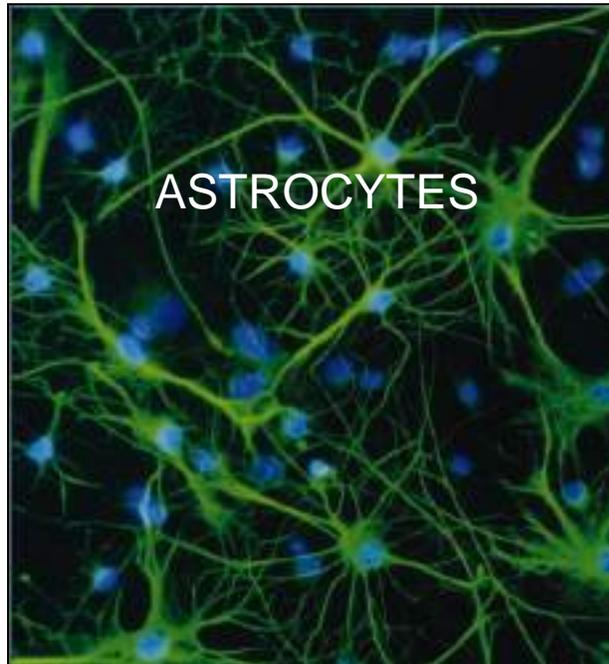
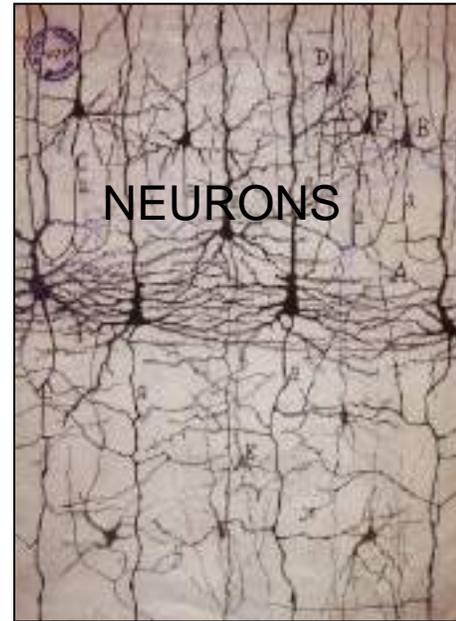
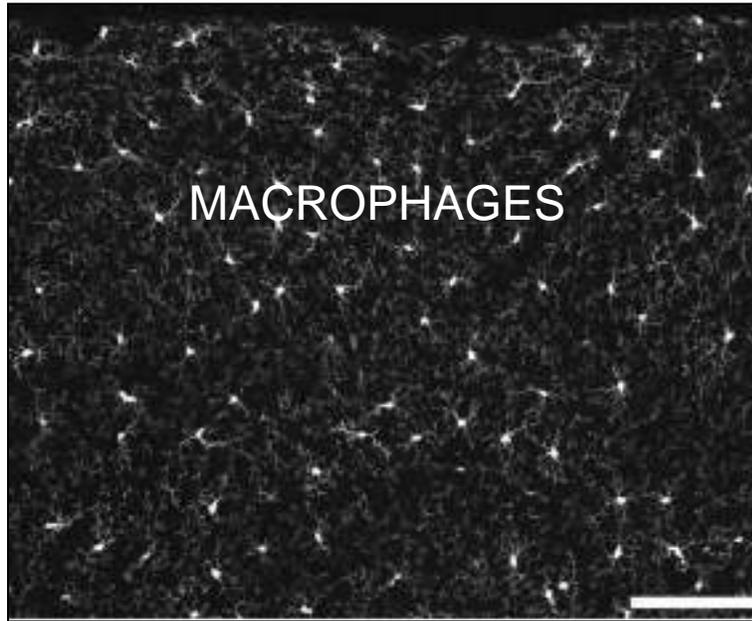


## Metabolic constraint imposes tradeoff between body size and number of brain neurons in human evolution

Fonseca-Azevedo K, Herculano-Houzel S

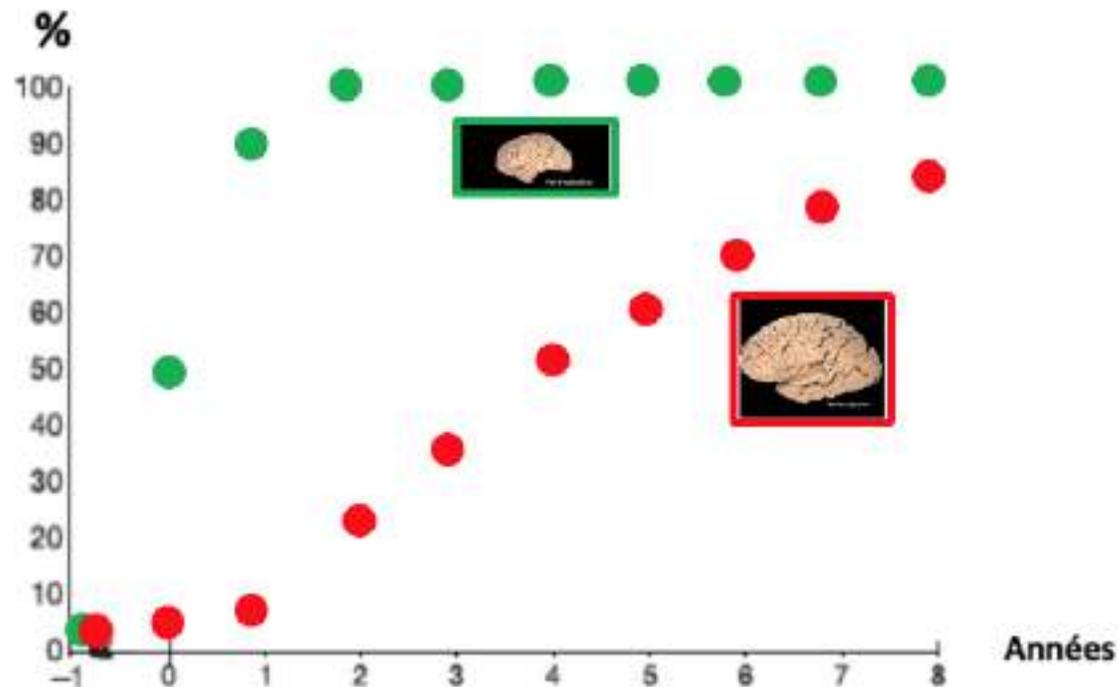
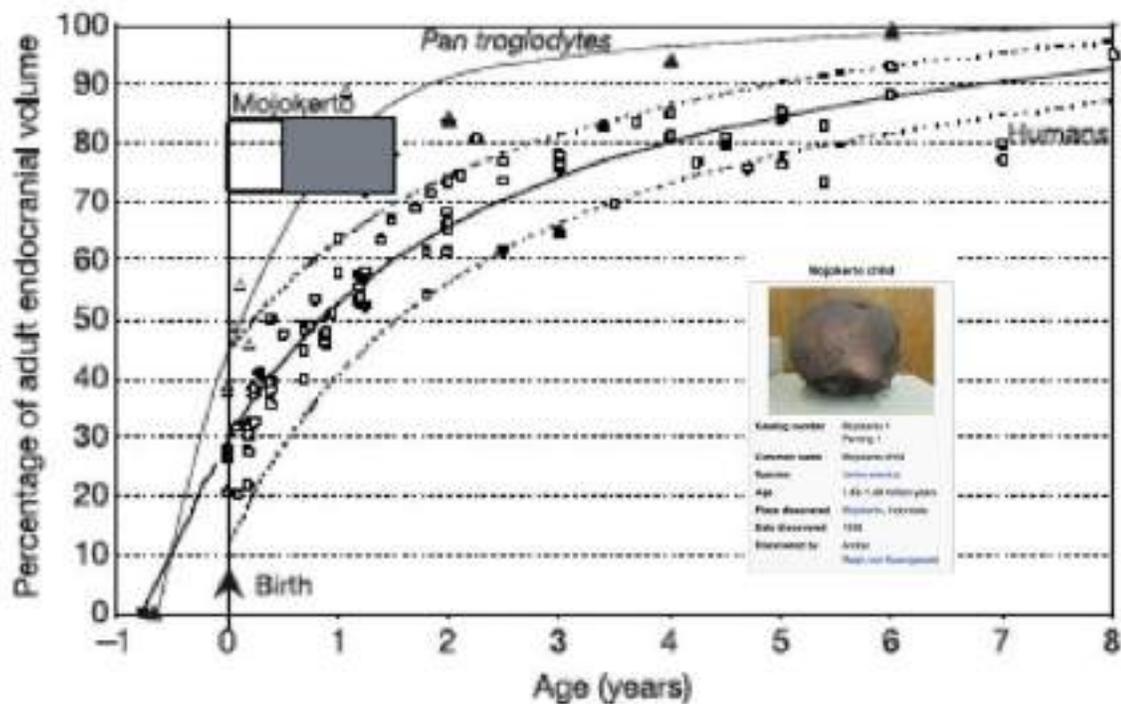
Proc Natl Acad Sci USA  
 2012 vol. 109 (45) pp. 18571-6

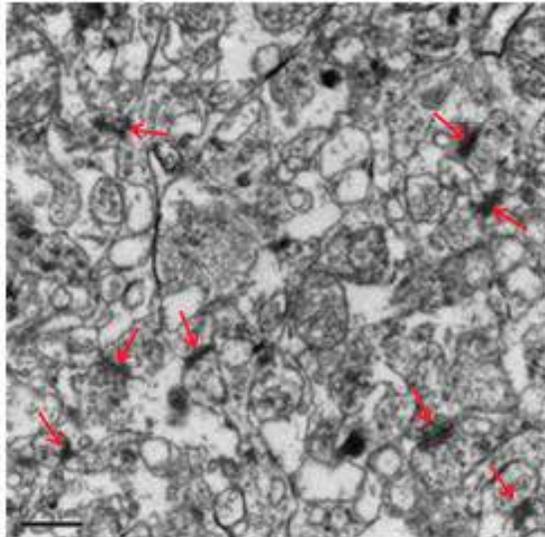




## Early brain growth in *Homo erectus* and implications for cognitive ability

H. Coqueugnot<sup>1</sup>, J.-J. Hublin<sup>2</sup>, F. Vellon<sup>3</sup>, F. Houët<sup>1</sup> & T. Jacob<sup>4</sup>



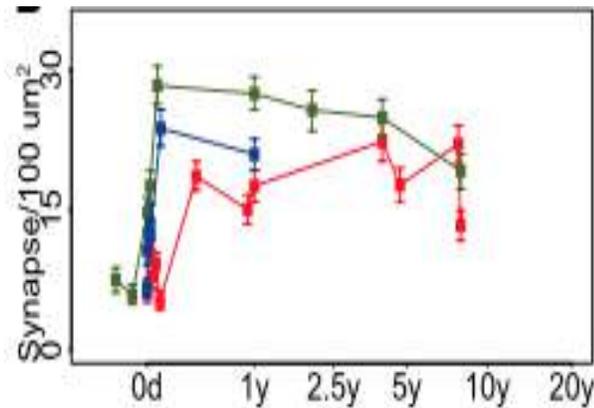
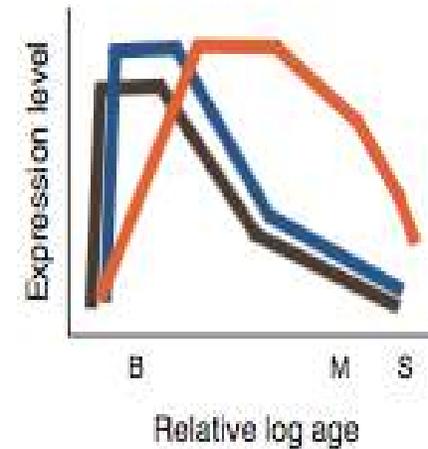


Extension of cortical synaptic development distinguishes humans from chimpanzees and macaques.

Xiling Liu et al.

Genome Research, 2012 vol. 22 (4) pp. 611-622

### Developmental remodeling

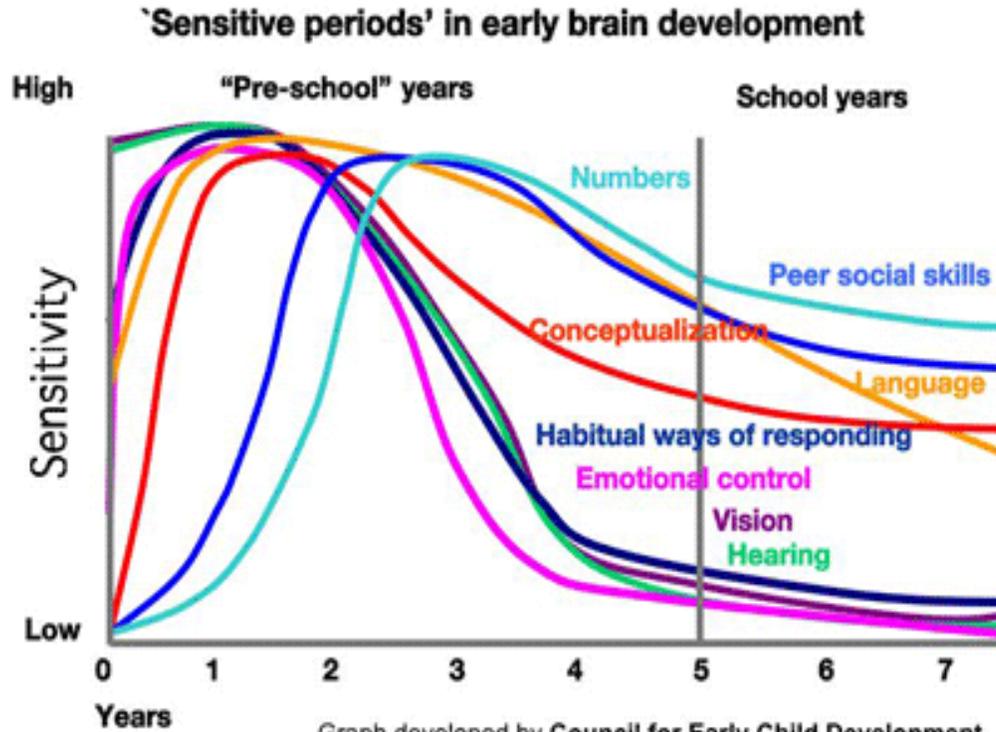


Human  
Chimp  
Macaque

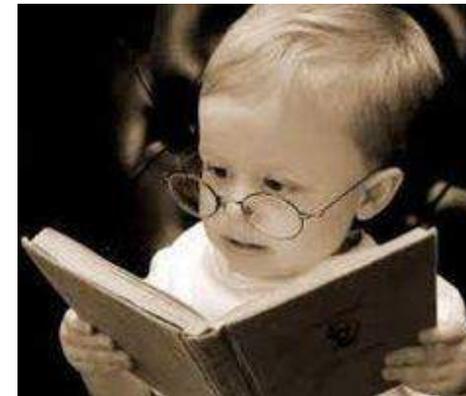
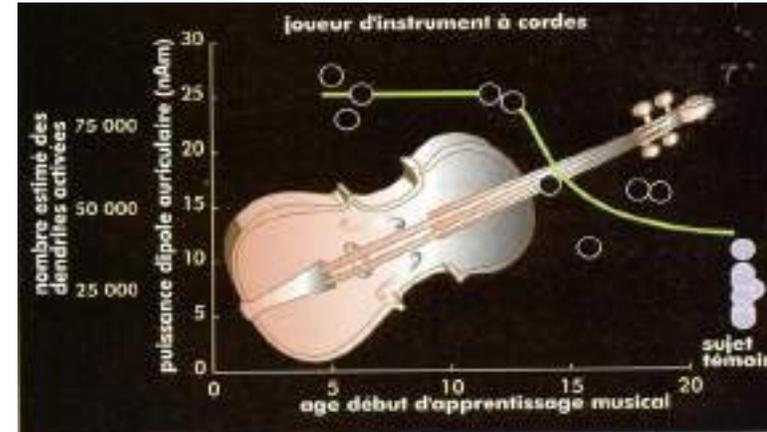


# Neuroplasticity

Ability to change neural pathways and synapses caused by changing behavior, environment, neural processes, thinking, emotions, as well as changes resulting from bodily injury

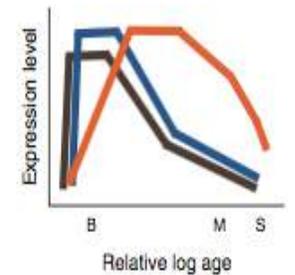


Graph developed by Council for Early Child Development (ref: Nash, 1997; Early Years Study, 1999; Shonkoff, 2000.)

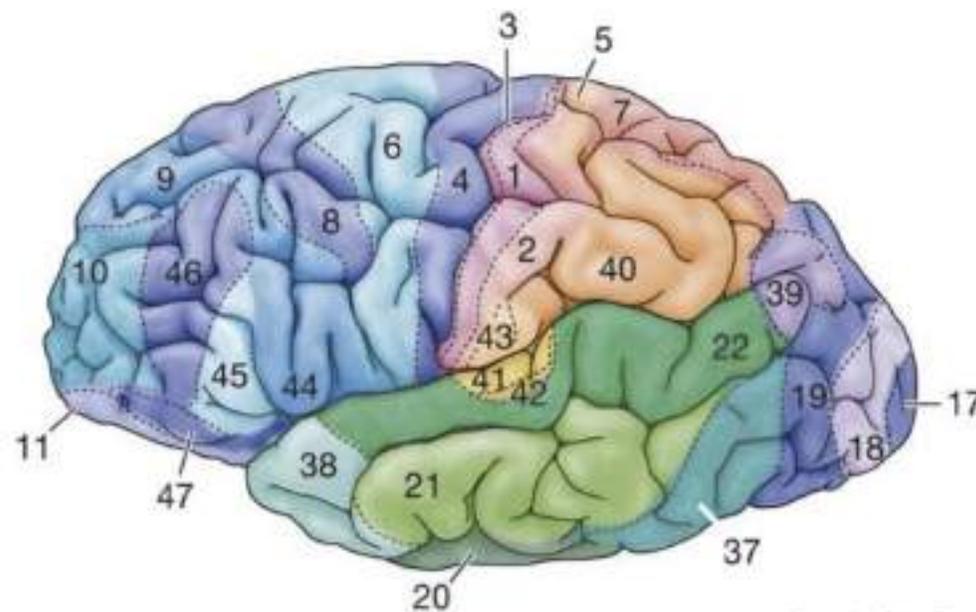
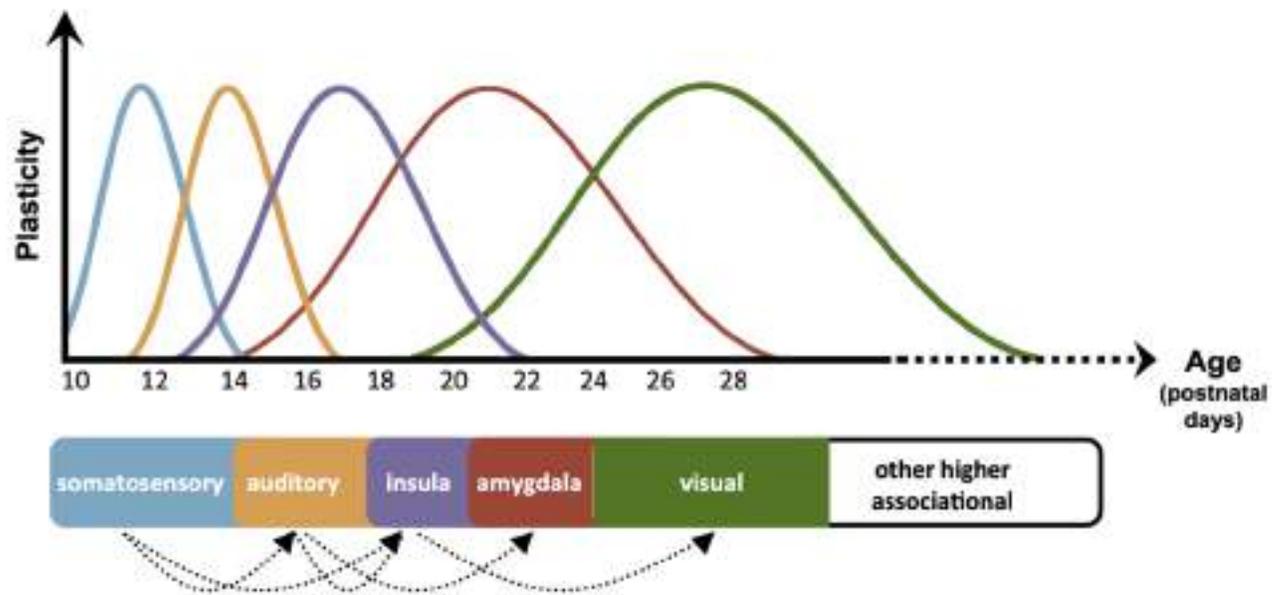


- Chimpanzee
- Macaque
- Human

Developmental remodeling

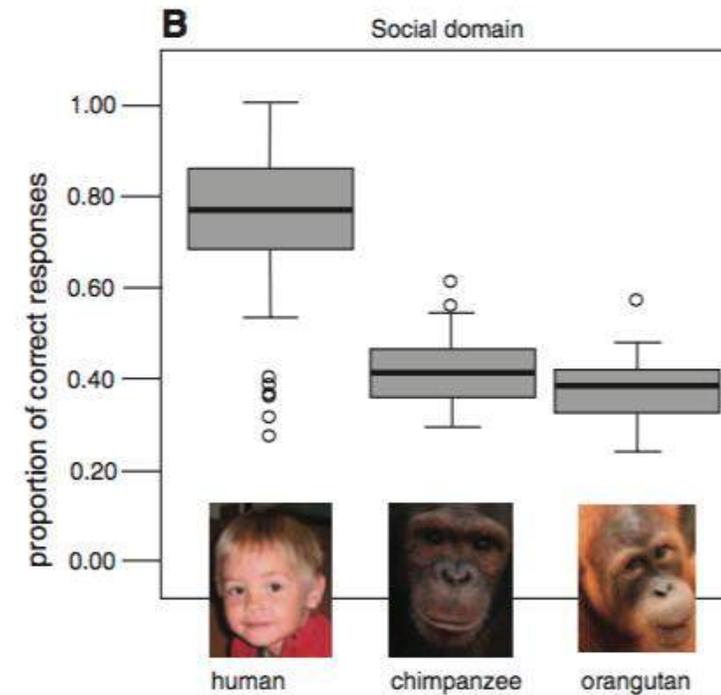
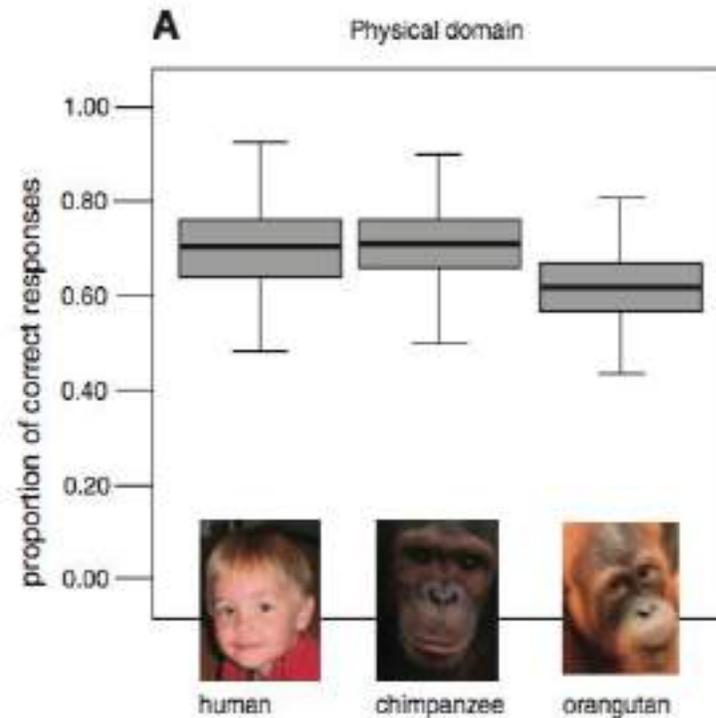


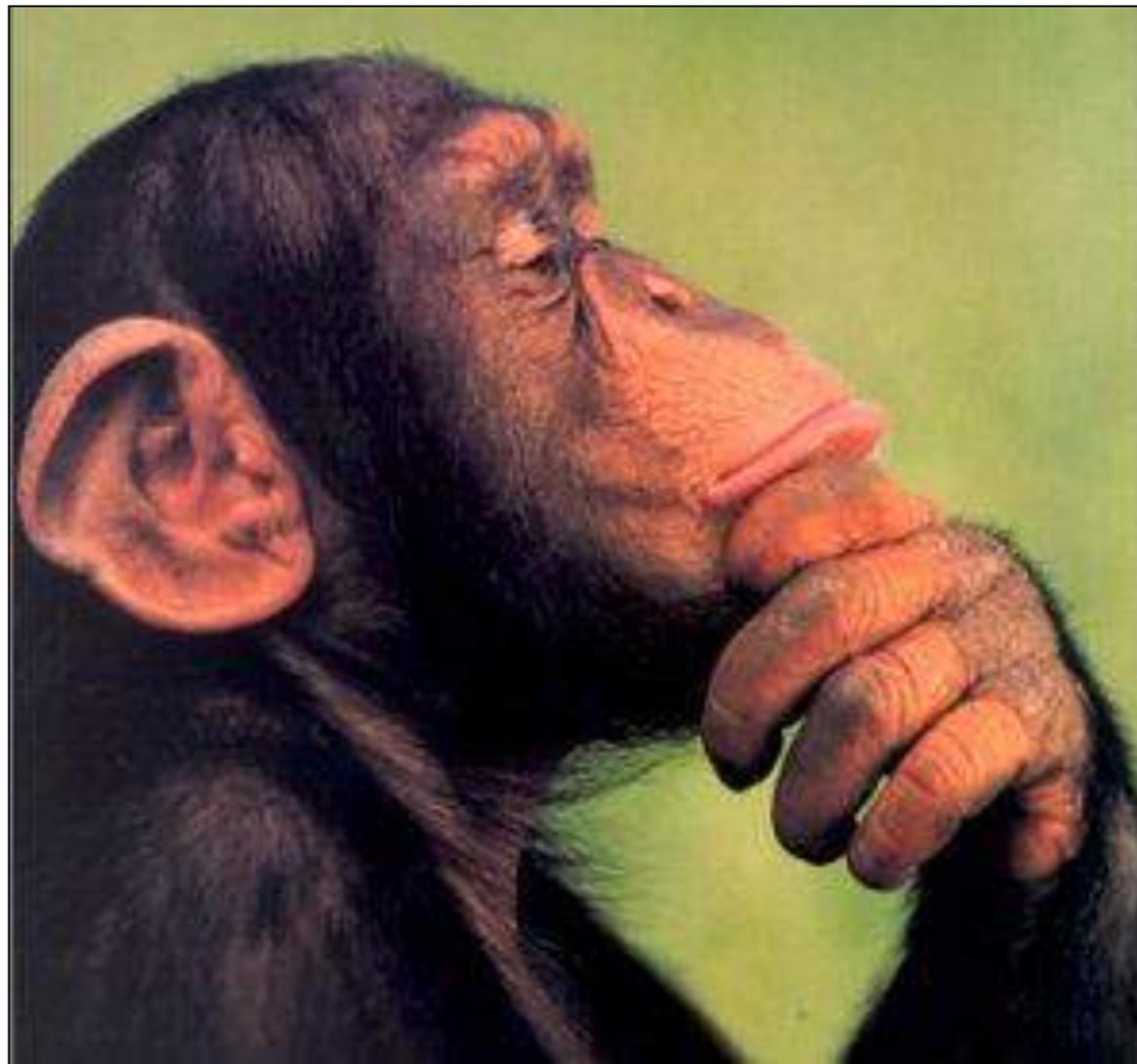
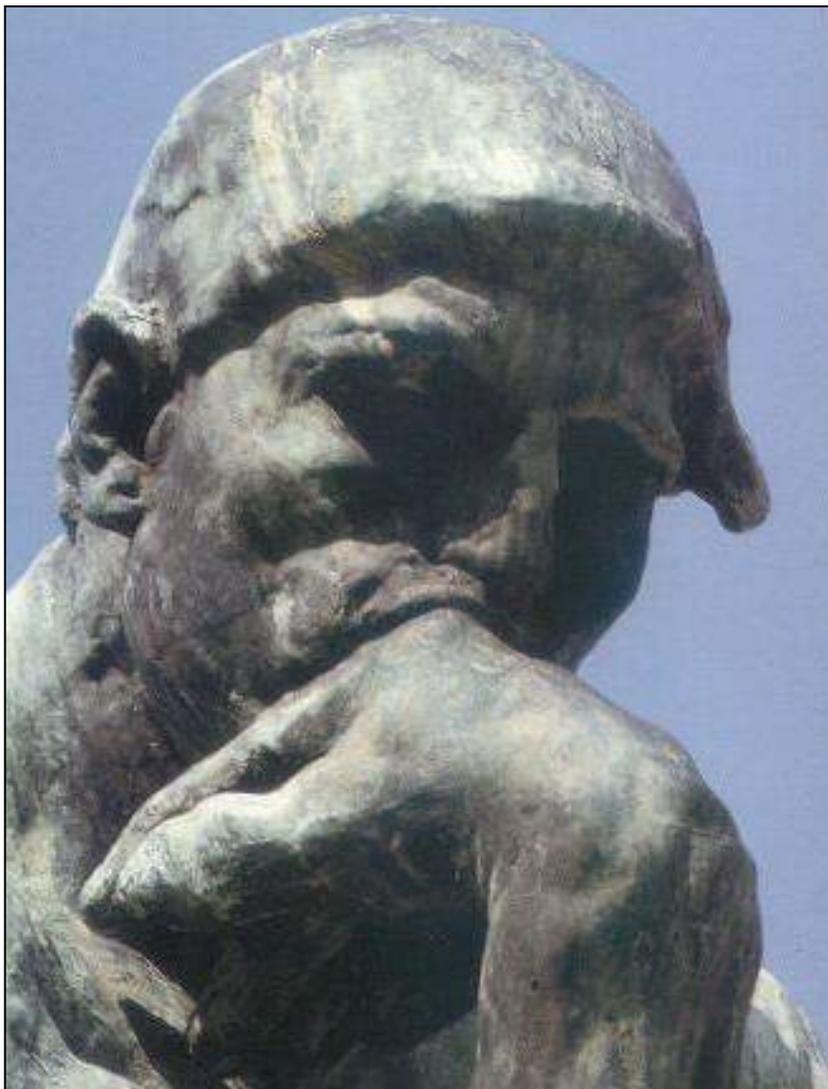
## Brain adapts to environment during heightened plasticity



# Humans Have Evolved Specialized Skills of Social Cognition: The Cultural Intelligence Hypothesis

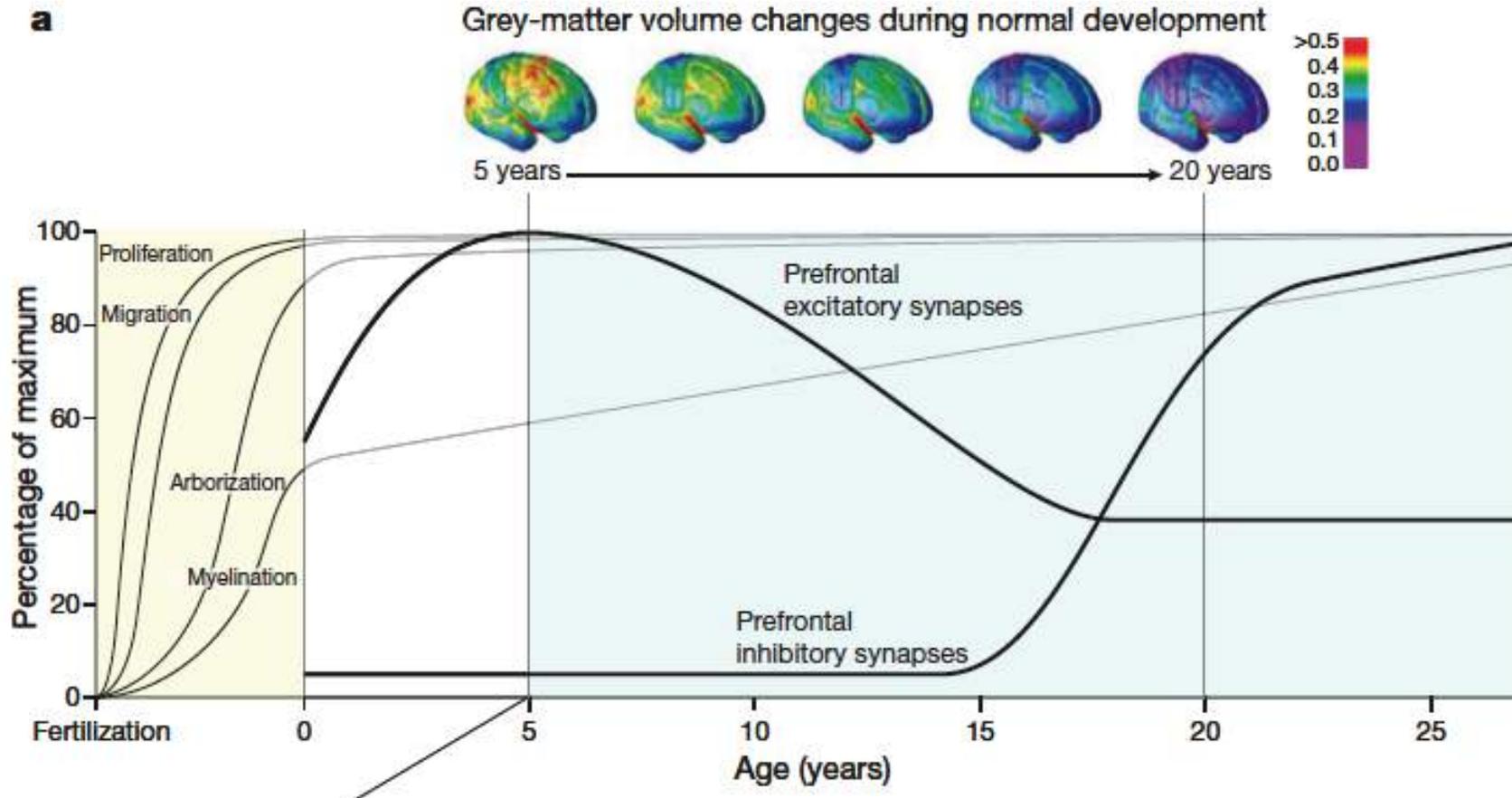
Esther Herrmann,<sup>1\*</sup> Josep Call,<sup>1</sup> María Victoria Hernández-Lloreda,<sup>2</sup>  
Brian Hare,<sup>3,3</sup> Michael Tomasello<sup>3</sup>





# Rethinking schizophrenia

Thomas S. Insel

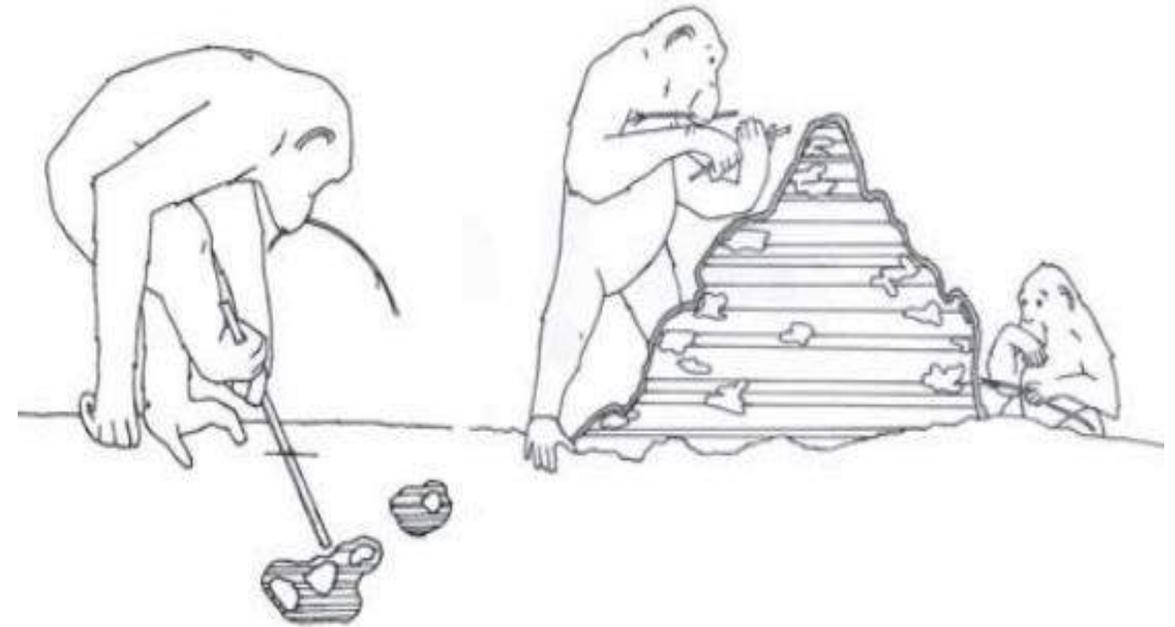
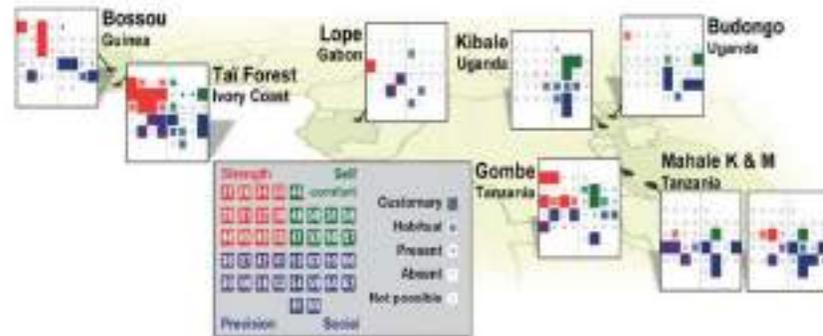


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# The second inheritance system of chimpanzees and humans

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Half a century of dedicated field research has brought us from ignorance of our closest relatives to the discovery that chimpanzee communities resemble human cultures in possessing suites of local traditions that uniquely identify them. The collaborative effort required to establish this picture parallels the one set up to sequence the chimpanzee genome, and has revealed a complex social inheritance system that complements the genetic picture we are now developing.



**Figure 1 | The cultures of wild chimpanzees.** Each chimpanzee community has its own unique array of traditions that together constitute the local 'culture'. 'Customary' acts are those typical in the community, 'habitual' ones are less common but consistent with social transmission, and 'absent' acts are those missing with no apparent straightforward environmental explanation. **Traditions are defined as behaviour patterns that are**

**customary or habitual in at least one site but absent elsewhere. Transmission is attributed to social learning on the basis of a complex of circumstantial evidence, ranging from intense observation by juveniles to distributions inconsistent with alternative explanations<sup>(1,2,3)</sup>. The numbers in cells refer to behaviour patterns in the catalogue of ref. 13, illustrated at <http://culture.st-and.ac.uk/chimp>.**

