

OPINION

Avian brains and a new understanding of vertebrate brain evolution

*The Avian Brain Nomenclature Consortium**

Abstract | We believe that names have a powerful influence on the experiments we do and the way in which we think. For this reason, and in the light of new evidence about the function and evolution of the vertebrate brain, an international consortium of neuroscientists has reconsidered the traditional, 100-year-old terminology that is used to describe the avian cerebrum. Our current understanding of the avian brain — in particular the neocortex-like cognitive functions of the avian pallium — requires a new terminology that better reflects these functions and the homologies between avian and mammalian brains.

One hundred years ago, Edinger, the father of comparative neuroanatomy, formulated a unified theory of brain evolution that formed the basis of a nomenclature that has been used to define the cerebral subdivisions of all vertebrates¹. This resulted in terms and associated concepts such as palaeostriatum, archistriatum, neostriatum and neocortex that are still in common use. According to this theory, the avian cerebrum is almost entirely composed of basal ganglia, the basal ganglia is involved in only instinctive behaviour, and the malleable behaviour that is thought to typify mammals exclusively requires the so-called neocortex. However, towards the end of the twentieth century, there accumulated a wealth of evidence that these viewpoints were incorrect. The avian cerebrum has a large pallial territory that performs functions similar to those of the mammalian cortex. Although the avian

pallium is nuclear, and the mammalian cortex is laminar in organization, the avian pallium supports cognitive abilities similar to, and for some species more advanced than, those of many mammals. To eliminate these misconceptions, an international forum of neuroscientists (BOX 1) has, for the first time in 100 years, developed new terminology that more accurately reflects our current understanding of the avian cerebrum and its homologies with mammals. This change in terminology is part of a new understanding of vertebrate brain evolution.

In this article, we summarize the traditional view of telencephalic evolution before reviewing more recent findings and insights. We then present the new nomenclature that has been developed by the Avian Brain Nomenclature Forum, and discuss its implications for our understanding of vertebrate brain evolution and its associated homologies.

The classical view

The classical view of telencephalic evolution, which is still prevalent in classrooms and textbooks, began in the late nineteenth and early twentieth centuries after the publication of *The Origin of Species* by Darwin². Inspired by Darwin's theory, between 1885 and 1908 Edinger formulated an influential evolution-based model of brain organization^{1,3,4}. Edinger and other early comparative neurobiologists combined Darwin's concept of 'evolution' with the nineteenth-century version of Aristotle's '*scala naturae*', which resulted in the view that evolution was progressive and unilinear⁵ — from fish, to amphibians, to

reptiles, to birds and mammals, to primates and, finally, to humans — ascending from 'lower' to 'higher' intelligence in a chronological series. They believed that the brains of extant vertebrates retained ancestral structures, and, therefore, that the origin of specific human brain subdivisions could be traced back in time by examining the brains of extant non-human vertebrates. In making such comparisons, they noted that the main divisions of the human CNS — the spinal cord, hindbrain, midbrain, thalamus, cerebellum and cerebrum or telencephalon — were present in all vertebrates (FIG. 1a). Edinger, however, noted that the internal organization of the telencephala showed the most pronounced differences between species. In mammals, the outer part of the telencephalon was found to have prominently layered grey matter (FIG. 1b, green) whereas the inner part had nuclear grey matter (FIG. 1b, purple). The inner part was located ventrally to the lateral ventricle. The outer part was more elaborate and folded in humans than in smaller mammals. In non-mammals, the outer and inner parts of the telencephala were mostly composed of nuclear grey matter, most of which was located ventrally to the lateral ventricle in reptiles and birds (FIG. 1b, purple).

On the basis of these considerations, Edinger proposed that telencephalic evolution occurred in progressive stages of increasing complexity and size, culminating with the human cerebrum. He suggested that the stages proceeded in a ventral-to-dorsal direction, with each new vertebrate group acquiring a more advanced cerebral subdivision, much as the earth's geological strata formed over time. He proposed that, first, there was the old brain, the palaeoencephalon (also called the basal ganglia or subpallium at the telencephalic base), which controlled instinctive behaviours, followed by the addition of a new brain, the neoencephalon (also called the pallium or mantle at the top of the telencephalon), which controlled learned and intelligent behaviours⁴. He, Ariëns Kappers

Box 1 | Avian Brain Nomenclature Consortium

Authors are ordered alphabetically in two groups: the first group, along with the first two and last two authors, are the core Avian Brain Nomenclature Forum Thinktank group; the second group are professors, postdoctoral fellows and students who also participated in the Avian Brain Nomenclature Forum. (For author affiliations see online [supplementary information S1](#) (box).)

Erich D. Jarvis, Onur Güntürkün, Laura Bruce, András Csillag, Harvey Karten, Wayne Kuenzel, Loreta Medina, George Paxinos, David J. Perkel, Toru Shimizu, Georg Striedter, J. Martin Wild, Gregory F. Ball, Jennifer Dugas-Ford, Sarah E. Durand, Gerald E. Hough, Scott Husband, Lubica Kubikova, Diane W. Lee, Claudio V. Mello, Alice Powers, Connie Siang, Tom V. Smulders, Kazuhiro Wada, Stephanie A. White, Keiko Yamamoto, Jing Yu, Anton Reiner and Ann B. Butler.

and others named the telencephalic subdivisions within each vertebrate group with the prefixes 'palaeo' (oldest), 'archi' (archaic) and 'neo' (new) to designate the presumed relative order of evolutionary appearance of each subdivision. In Greek, 'archi' means the oldest, the first, or the most primitive, whereas 'palaeo' means ancient, primitive or old, but not necessarily the oldest. Both Edinger and Ariëns Kappers misinterpreted the meaning of these prefixes and reversed them, naming structures with 'palaeo-' to indicate the oldest or first and 'archi-' to indicate old. They added to these prefixes the root word 'striatum' for the presumed palaeoencephalic subdivisions and 'pallium' or 'cortex' for the presumed neoencephalic subdivisions^{1,4,6-8}. The term 'striatum' was used because a large part of the basal ganglia (palaeoencephalon) in mammals, now commonly called the caudate-putamen, has fibre bundles coursing through it that give it a striated appearance.

The classical view that became dominant was that the primordial telencephalon of fishes had a relatively small pallium and a larger subpallium, both of which were entirely devoted to olfactory information processing. The fish subpallium was named 'palaeostriatum' (old striatum), and was thought to be the antecedent of the human globus pallidus (FIG. 1b). Amphibians were thought to have evolved an 'archistriatum' (archaic striatum) above the palaeostriatum, which was proposed to be the antecedent of the human amygdala. Reptiles were thought to have evolved a 'neostriatum' (new striatum) above the archistriatum, which was proposed to be the antecedent of the human caudate and putamen. The palaeostriatum of reptiles was also thought to have elaborated into an older part (primitivum) and a newer part (augmentatum), both of which were considered homologous to the human globus pallidus. Following this, birds were thought to have evolved a large additional basal ganglia subdivision, the 'hyperstriatum' (hypertrophied striatum), which was considered to be unique to birds⁹.

The fish pallium was named 'palaeocortex', and was proposed to be the antecedent of the human olfactory cortex. Reptiles were thought to have evolved an 'archicortex', also thought to be olfactory and primitive, that was said to be the antecedent of the human hippocampus. Birds were thought not to have evolved any further pallial regions. By contrast, mammals were thought to have evolved the latest and greatest achievement, a 'neocortex', from the palaeocortex and/or archicortex⁶. The archicortex and/or palaeocortex, with their 2–3 cell layers, were assumed to be primitive; the neocortex, with its 6 layers, was assumed to be more recently evolved and a substrate for more sophisticated behaviour.

There were dissenting voices to the classical view¹⁰⁻¹². Some of its proponents also made partial or tentative retractions^{13,14}. However, alternative views were not widely embraced. Instead, the classical view was codified in the important 1936 comparative neuroanatomy text by Ariëns Kappers, Huber and Crosby¹⁴ and became pervasive throughout neuroscience.

A new view of the subpallium Substantive challenges to the classical view of the subpallium began in the 1960s and 1970s with the advent of new methods for determining both nervous system connectivity and the anatomical profiles of gene products⁵. These studies found that, in mammals, acetylcholinesterase enzymatic activity was enriched in the neostriatum¹⁵. In birds, high acetylcholinesterase activity was found only in the palaeostriatum augmentatum and associated lobus parolfactorius (LPO)^{15,16} (the LPO was considered to be part of the palaeostriatum augmentatum by Ariëns Kappers *et al.*¹⁴, but was named as a separate region by Karten and Hodos¹⁷). Other studies found that the mammalian neostriatum was highly enriched with dopaminergic terminals, which originated from midbrain neurons in the substantia nigra pars compacta¹⁸. In birds, again, only the palaeostriatum augmentatum and LPO were enriched with dopaminergic terminals, and

this input originated from neurons in the midbrain^{19,20}. During the following decades, using new methods in double-label immunohistochemistry and tract tracing, the mammalian neostriatum was found to be enriched in two types of neuron: those containing the neuropeptide substance P (SP), which project to the internal part of the globus pallidus and substantia nigra, and those containing the neuropeptide enkephalin (ENK), which project to the external part of the globus pallidus²¹⁻²⁴. In birds, SP and ENK neurons are enriched in the palaeostriatum augmentatum (including the LPO)^{24,25}, and, like the equivalent neurons in mammals, project to different cell types within the adjacent avian palaeostriatum primitivum. In both birds and mammals the SP neurons seem to be involved in promoting planned movement, whereas the ENK neurons seem to have a role in inhibiting unwanted movement. Further functional studies revealed that both the mammalian neostriatum and avian palaeostriatum augmentatum (including the LPO) participate not only in instinctive behaviour and movement, but also in motor learning^{26,27}.

These apparent relationships between the subpallia of mammals and birds have been supported by molecular embryology studies^{24,28-31}. The developing subpallium in birds and mammals consists of two separate histogenetic zones that express different sets of transcription factors: a dorsal zone, which, in mammals, corresponds to the lateral ganglionic eminence and selectively expresses the transcription factors *DLX1* and *DLX2* but not *NKX2.1*; and a ventral zone, which, in mammals, corresponds to the medial ganglionic eminence and selectively expresses all three transcription factors. In mammals, the lateral ganglionic eminence gives rise to the dorsal striatum (neostriatum) and the ventral striatum (nucleus accumbens and part of olfactory tubercle). The homologous developing territory in birds gives rise to the structures that were previously called the palaeostriatum augmentatum (including the LPO) and the olfactory tubercle. The medial ganglionic eminence in mammals gives rise to various pallidal cell groups, including the dorsal pallidum (the globus pallidus) and the ventral pallidum. The homologous developing territory in birds gives rise to the structures that were called the palaeostriatum primitivum and ventral palaeostriatum. These avian and mammalian striatal and pallidal relationships are further supported by studies of the comparative expression patterns of more than 30 other genes in adult birds and mammals³²⁻³⁶. Similar striatal and pallidal territories have been found in the so-called palaeostriatal regions of reptiles³⁷⁻⁴⁰.

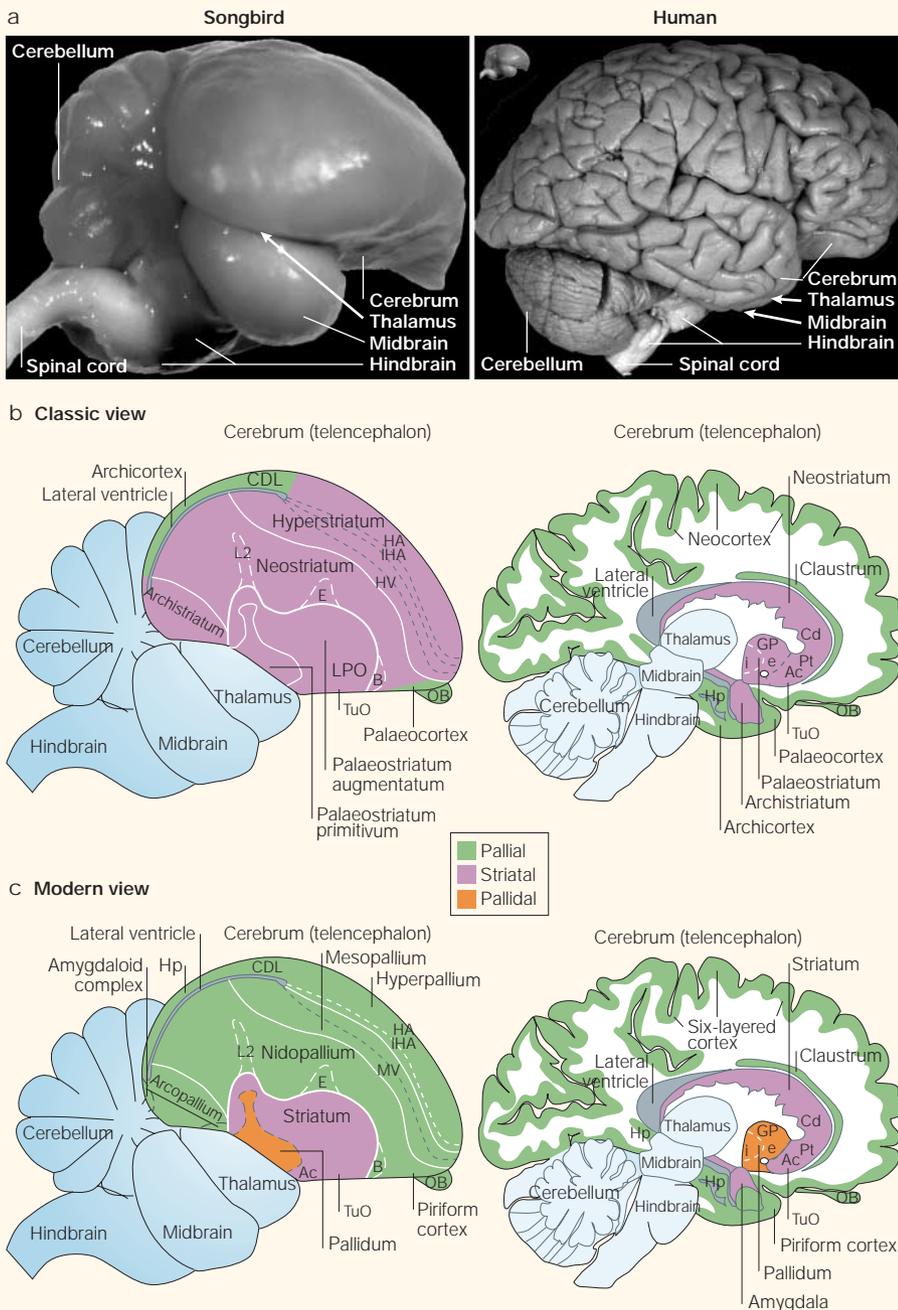


Figure 1 | Avian and mammalian brain relationships. **a** | Side view of a songbird (zebra finch) and human brain to represent avian and mammalian species. In this view, the songbird cerebrum covers the thalamus; the human cerebrum covers the thalamus and midbrain. Inset (left) next to the human brain is the zebra finch brain to the same scale. Human brain image reproduced, with permission, courtesy of John W. Sundsten, Digital Anatomist Project. **b** | Classic view of avian and mammalian brain relationships. Although past authors had different opinions about which brain regions are pallium versus subpallium, we have coloured individual brain regions according to the meaning of the names given to those brain regions. Ac, accumbens; B, nucleus basalis; Cd, caudate nucleus; CDL, dorsal lateral corticoid area; E, ectostriatum; GP, globus pallidus (i, internal segment; e, external segment); HA, hyperstriatum accessorium; HV, hyperstriatum ventrale; IHA, interstitial hyperstriatum accessorium; L2, field L2; LPO, lobus parolfactorius; OB, olfactory bulb; Pt, putamen; TuO, olfactory tubercle. **c** | Modern consensus view of avian and mammalian brain relationships according to the conclusions of the Avian Brain Nomenclature Forum. Solid white lines are lamina (cell-sparse zones separating brain subdivisions). Large white areas in the human cerebrum are axon pathways called white matter. Dashed grey lines divide regions that differ by cell density or cell size; dashed white lines separate primary grey neuron populations from adjacent regions. Abbreviations where different from **b**: E, entopallium; B, basorostralis; HA, hyperpallium apicale; Hp, hippocampus; IHA, interstitial hyperpallium apicale; MV, mesopallium ventrale.

Together, these studies indicate that the avian palaeostriatum augmentatum is homologous to the mammalian neostriatum and that the avian palaeostriatum primitivum is homologous to the mammalian globus pallidus.

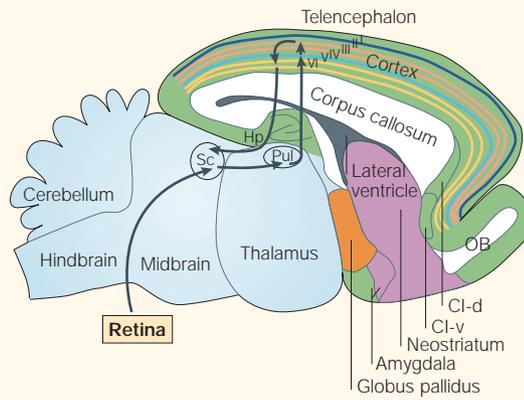
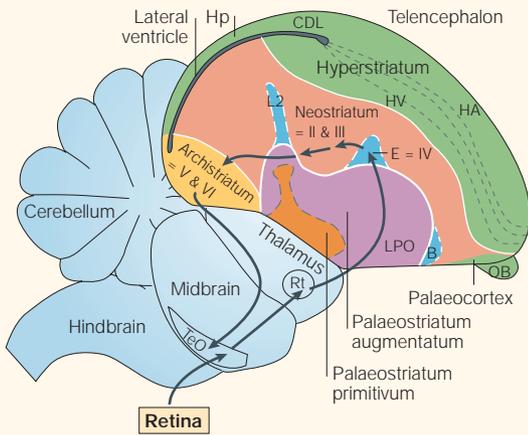
A new view of the pallium

With these challenges to the classical view of the subpallial relationships among birds, reptiles and mammals came challenges to the classical view of the relationships among their pallia. The mammalian pallium includes the areas known as palaeocortex, archicortex and neocortex; and has been said, more recently, to include both the claustrum and lateral parts of the amygdala^{28,41,42} (FIG. 1c; Holmgren¹¹ originally proposed that the claustrum and part of the amygdala were pallial, but this view was largely ignored at the time). In birds, the finding that the structures that had been called hyperstriatum, neostriatum and archistriatum were neither striatum nor pallidum raised the question of which telencephalic sector these regions did represent. The results that were needed to answer this question also began to appear in the mid-1960s from pathway tracing^{16,43–46} and behavioural studies^{47–51}. These studies found that the so-called avian neostriatum and hyperstriatum receive visual, auditory and somatosensory input from the thalamus, as does the mammalian neocortex. These avian brain regions also carry out the same type of sensory information processing as is performed by the mammalian neocortex. The so-called avian hyperstriatum accessorium and the archistriatum give rise to important descending projections to the premotor and motor neurons of the brainstem and spinal cord, similar to those of the mammalian cortico-bulbar and cortico-spinal pathways^{43,52–54}. Finally, like the mammalian neocortex, these avian brain regions carry out crucial roles in motor control and sensorimotor learning^{55–67}.

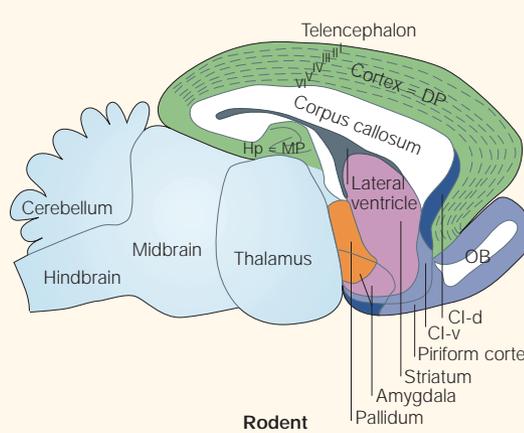
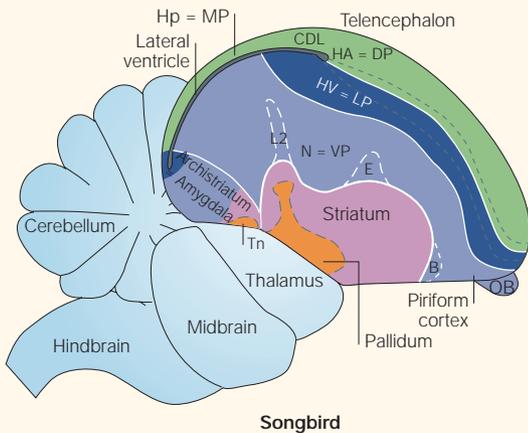
The apparent pallial relationships between these mammalian and avian brain regions were also supported by molecular embryology studies^{28,68,69}. During development, both the avian hyperstriatum and neostriatum and the mammalian pallium express the pallium-specific transcription factors *EMX1*, *PAX6* and *TBR1*. The developmental data led to uncertainties about how much of the archistriatum is pallial^{28,30}. However, comparisons of the expression of the brain-derived neurotrophic factor (*BDNF*) and the glutamate receptor *mGluR2* in adult birds and mammals indicated that the entire avian archistriatum, as defined in brain atlases^{17,70}, expresses these pallium-specific mRNAs^{34,36}. Further studies of the comparative expression patterns of other glutamate receptors in adult birds and mammals³⁶

Box 2 | Working hypotheses on avian and mammalian pallial homologies.

a Nuclear-to-layered hypothesis



b Nuclear-to-claustrum/amygdala hypothesis



An example of a nuclear-to-layered hypothesis is shown in panel a. The connectivity of tectofugal visual pathways in avian (left) and mammalian (right) brains is shown. The hypothesis illustrated is that of Karten⁷¹. Colour-coding indicates proposed homologies between birds and mammals. An example of a nuclear-to-claustrum/amygdala nuclei hypothesis is shown in panel b. The hypothesis illustrated is that of Puelles *et al.*²⁸.

I–VI, cortical layers I–VI; B, nucleus basalis; CDL, dorsal lateral corticoid area; Cl-d, claustrum, dorsal part; Cl-v, claustrum, ventral part; DP, dorsal pallium; E, ectostriatum; HA, hyperstriatum accessorium; Hp, hippocampus; HV, hyperstriatum ventrale; L2, field L2; LP, lateral pallium; LPO, lobus parolfactorius; MP, medial pallium; N, Neostriatum; OB, olfactory bulb; Pul, pulvinar nucleus; Rt, nucleus rotundus; Sc, superior colliculus; TeO, optic tectum; Tn, nucleus taenia; VP, ventral pallium.

support these conclusions. Together, these studies indicate that the avian hyperstriatum, neostriatum, and archistriatum might be homologous to mammalian pallial regions.

This developing view was accompanied by several new proposals about one-to-one homologies between specific avian and mammalian pallial subdivisions. We will consider these in two groups — nuclear-to-layered hypotheses and nuclear-to-claustrum/amygdala nuclei hypotheses.

Nuclear-to-layered hypotheses. First proposed by Karten^{16,71}, nuclear-to-layered hypotheses (BOX 2) propose that the similarities in connectivity between the so-called hyperstriatum, neostriatum and archistriatum of birds and the neocortex of mammals stem from a common origin of these structures — that is, they are homologous. Karten proposed that the common ancestor of birds, reptiles and mammals possessed a nuclear pallium that was transformed into a laminar pallium early in

the mammalian lineage, maintaining the connectivity of the ancestral nuclear network. In this regard, he argued that the avian pallium is divided into three groups of serially connected neuron types — thalamorecipient neurons (field L2, ectostriatum and basalis), pallio-pallial neurons (neostriatum) and extratelencephalic projection neurons (archistriatum), with cell types and interconnectivity that resemble those of mammalian cortical layers IV, II–III and V–VI, respectively (BOX 2). Similar arguments were later made for the avian upper hyperstriatum (also known as the Wulst), which also has serially connected neuron types that resemble those found in the mammalian neocortex⁶². In this hypothesis, avian L2 neurons are homologous to layer IV neurons of mammalian primary auditory cortex, basalis neurons to layer IV of primary somatosensory cortex, ectostriatal neurons to layer IV of extrastriate visual cortex, and the interstitial hyperstriatum accessorium to layer IV of striate visual cortex. In support of this

hypothesis, gene expression studies^{36,72,73} have shown that avian thalamorecipient nuclear fields (L2, ectostriatum, basalis and interstitial hyperstriatum accessorium) and the mammalian thalamorecipient layer IV of neocortex selectively express some of the same genes (the steroid transcription factor *ROR-β* and the potassium channel *EAG2*) and express a low level of others (the activity-dependent transcription factor *ZENK* and the AMPA (α-amino-3-hydroxy-5-methyl-4-isoxazole propionic acid) glutamate receptor subunit *GluR1*). Avian extratelencephalic projection neurons (in the archistriatum, but not in the hyperstriatum accessorium) and mammalian extratelencephalic projection neurons (layer V neurons of neocortex) both show selective expression of the transcription factor *ER81*. So, although the avian pallium is not organized cytoarchitectonically into layers, its nuclear subdivisions bear marked similarities in connectivity and molecular profile to different layers of the mammalian neocortex.

Nuclear-to-claustrum/amygdala hypotheses.

These hypotheses (BOX 2) provide a different interpretation of mammalian homologies with the avian ventral hyperstriatum, neostriatum and archistriatum, known collectively as the dorsal ventricular ridge (DVR). In an early proposal, Bruce and Neary⁷⁴ proposed that the avian DVR represents an elaboration of parts of the mammalian amygdala. Subsequently, Striedter⁷⁵ proposed that the avian DVR represents an elaboration of the mammalian amygdala and claustrum, and that the connectivity that the DVR shares with the neocortex evolved independently. Support for this view was based on several facts: both the avian DVR and mammalian claustrum/amygdala are nuclear in organization⁷⁵; both the avian DVR and part of the mammalian amygdala have similar connections^{74,76,77}; and both have conserved developmental expression patterns of regulatory genes that have important roles in brain regionalization and morphogenesis^{28,68}. In the most detailed gene expression study, Puelles *et al.*²⁸ proposed that the common topographic expression patterns of the transcription factors *EMX1* and *PAX6* in the avian hyperstriatum ventrale and in the mammalian dorsal claustrum and basolateral amygdala indicate that these structures both arose from the lateral pallium (BOX 2). They argued that the absence of *EMX1* but the presence of other pallial genes in the avian neostriatum and in the mammalian ventral claustrum and lateral anterior amygdala indicate that these structures commonly arose from the ventral pallium. They further proposed that the avian archistriatum and mammalian amygdala consist of subpallial parts derived from striatal and pallidal cell groups, and, by this association, that the avian archistriatum is homologous to the mammalian amygdala, as originally proposed by Edinger¹.

Both the above hypotheses have their limitations. For the nuclear-to-layered hypotheses, developmental studies have not been conducted to investigate whether the three types of serially connected neuron in birds arise from cell types similar to those that give rise to the cortical layers in mammals. Furthermore, not all gene expression patterns support one-to-one molecular relationships between avian pallial subdivisions and mammalian cortical layers⁷⁸. In addition, not all findings support the nuclear-to-claustrum/amygdala hypothesis. Although initial studies²⁸ reported that this hypothesis was supported by the lack of pallial *EMX1* expression in the avian neostriatum (so-called ventral pallium) and the

mammalian ventral claustrum and lateral anterior amygdala, recent fate mapping showed that the ventral claustrum contains dispersed *EMX1*-expressing cells and the lateral amygdala contains many *EMX1*-expressing cells⁷⁹. The antiquity of the claustrum has also been debated. One study⁸⁰ reported that monotremes (platypus and echidnas) lack a claustrum, whereas a later study⁸¹ reported that echidnas, but not platypus, have a rudimentary claustrum located more ventrally in the white matter relative to the location of the claustrum in other mammals. So, it is possible that some monotreme groups have lost the claustrum; that the echidna has independently evolved it or another deep cortical derivative; or that the claustrum is not an ancestral mammalian trait. Further investigation is required.

A new nomenclature

Despite an extensive revision of our understanding of telencephalic evolution, the common nomenclature used for the avian telencephalon has, until 2004 (REF. 82), retained all of the classical evolution- and *scala naturae*-based terminology. For this reason, findings in 'birdbrains' have been habitually misinterpreted by neuroscientists studying non-avian brains as pertaining to the basal ganglia or as largely irrelevant to mammals. To rectify this problem, an international consortium of specialists in avian, mammalian, reptilian and fish neurobiology — the Avian Brain Nomenclature Consortium (BOX 1) — assembled with the goal of revising the terminology for the avian brain. Through online discussions, an Avian Brain Nomenclature Exchange web site, various meetings held over a period of 6 years and an Avian Brain Nomenclature Forum held at Duke University, North Carolina⁸³, the group developed a new terminology that represents the current understanding of avian telencephalic organization and its homologies with mammals⁸² (FIG. 1c). On the basis of the evidence summarized above, we concluded that the avian telencephalon is organized into three main, developmentally distinct domains that are homologous in fish, amphibians, reptiles, birds and mammals: pallial, striatal and pallidal domains (FIG. 1c). We renamed the subdivisions within each of these domains in birds with homology-based terms or roots that allow reference to named regions in mammals, and eliminated all phylogeny-based prefixes (palaeo-, archi- and neo-) that erroneously implied the relative age of each subdivision.

The striatal and pallidal domains. We renamed the avian palaeostriatum augmentatum and LPO as the lateral and medial parts of the avian dorsal striatum, and identified a nucleus accumbens and olfactory tubercle as parts of the avian ventral striatum (FIG. 1c). We renamed the region that includes the palaeostriatum primitivum and ventral palaeostriatum as the 'pallidum' (FIG. 1c). Like the mammalian pallidum, the avian pallidum has a sparse distribution of cells²⁴, giving the region its pale appearance and, therefore, its name. The dorsal region of the avian pallidum was found to be homologous to the mammalian globus pallidus and named as such, whereas the ventral part was determined to be homologous to the mammalian ventral pallidum. The dorsal pallidum, however, differs between mammals and birds. In mammals, it consists of two segments with distinct connectivity — the internal and external globus pallidus — whereas in birds, neurons with both phenotypes are intermingled²⁵ (FIG. 1c).

The pallial domain. We concluded that the avian pallium is organized into four main subdivisions instead of three striatal subdivisions (hyperstriatum, neostriatum and archistriatum)⁷ and renamed them hyperpallium (hypertrophied pallium; upper part of old hyperstriatum), mesopallium (middle pallium; lower part of old hyperstriatum), nidopallium (nest pallium; old neostriatum) and arcopallium (arched pallium; most of old archistriatum) (FIG. 1c). We concluded that several neuronal populations adjacent to the arcopallium and the posterior part of what had been regarded as archistriatum are homologous to pallial and subpallial regions of the mammalian amygdala, and renamed them as members of the amygdaloid complex. Other regions that were widely recognized to be homologous among vertebrates — the hippocampus, olfactory (piriform) cortex and olfactory bulb — did not require name changes. After extensive evaluation of the various one-to-one homology hypotheses of the avian and mammalian pallia^{28,44,68,74,75,77,84} (BOX 2), we concluded that the evidence is not strong enough for any specific proposed homologies to be incorporated into a new pallial terminology. However, we recognize that this is an area of active research and debate, and designed the new terminology to be compatible with the adoption of any one-to-one homology hypothesis should future evidence be more convincing.

A new view of telencephalic evolution
With this new understanding of the avian telencephalic organization and its homologies

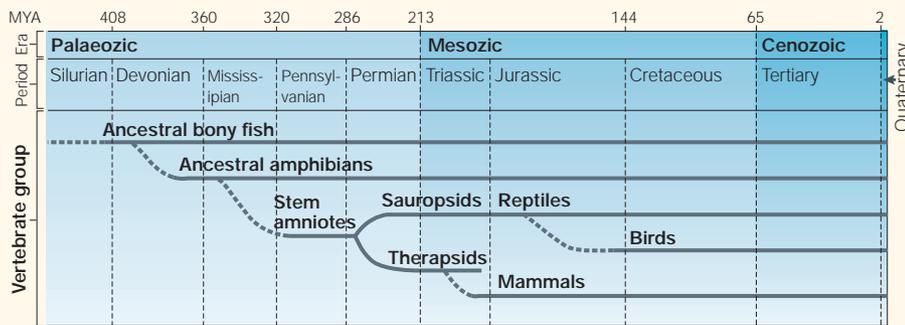


Figure 2 | **Simplified modern view of vertebrate evolution.** The diagram begins with the fish group that contains the most recent ancestors of land vertebrates. This differs from the classic view in that instead of giving rise to reptiles, ancestral amphibians are thought to have given rise to stem amniotes. Stem amniotes then split into at least two groups: the sauropsids, which gave rise to all modern reptiles as we know them today; and the therapsids, which, through a series of now-extinct intermediate forms, evolved into mammals. Many sauropsids (reptiles) are currently living. Solid horizontal lines indicate temporal fossil evidence. Dashed lines indicate proposed ancestral links based on other types of data. Based on REFS 85,86.

with that of mammals, we can generate more informed hypotheses and conclusions about telencephalic evolution in vertebrates. It is now apparent that the organization of the true basal ganglia among birds, mammals and other vertebrates (that is, distinct nuclear striatal and pallidal domains with more dopaminergic input into the striatal domain²⁴) is quite conserved. By contrast, the organization of the pallial domains of these groups is more varied. The avian hyperpallium has a unique organization that has so far been found only in birds⁶⁹. This consists of semi-layered subdivisions, and might have evolved more recently than the mammalian six-layered cortex, as birds evolved ~50–100 million years after mammals^{85,86} (FIG. 2). The DVR (which, in birds, contains the mesopallium, nidopallium and arcopallium) is a nuclear, grey matter formation that is unique to birds and reptiles. The six-layered cortex is unique to mammals, and, as all the main groups of living mammals (monotremes, marsupials and placentals) have a six-layered cortex⁸⁷, it was presumably inherited from their common therapsid ancestor more than 200 million years ago (FIG. 2). Furthermore, new findings indicate that mammals did not arise from reptiles, but from therapsids, and that the last common ancestor of the reptile and mammal lineages was the stem amniotes⁸⁶. As all non-mammalian therapsids are now extinct (FIG. 2), it is difficult to trace from stem amniotes to mammals the evolutionary history of mammalian telencephalic organization — layered, nuclear or otherwise. Therefore, the reptilian nuclear pallial organization cannot be assumed to represent the ancestral condition for mammals.

Further, it is now known that evolution is not invariably progressive or linear, so there is no basis for the view that more recently evolved species or structures are more

advanced. In support of this conclusion, we now know that the telencephala of fishes are not devoted mainly to olfactory function, as the olfactory area represents only a limited portion of the fish pallium⁸⁸. In addition, fishes have a hippocampus (archicortex), and the main function of the hippocampus in fish, reptiles, birds and mammals alike is not olfaction, but memory formation and spatial mapping^{89,90}.

So, as for birds, it might be best to abandon the use of the terms archicortex, palaeocortex, archistriatum, palaeostriatum and neostriatum for mammals and other vertebrates, in favour of the alternatives — hippocampus or hippocampal cortex, piriform cortex, amygdala, striatum and pallidum. However, alternative terms for ‘neocortex’, such as ‘isocortex’, have not been universally accepted. ‘Neocortex’ would be appropriate if taken to refer to the uniqueness of this cortical structure among vertebrates. However, ‘neocortex’ should not be taken to mean that it is the only unique form of pallial organization, that it evolved out of a palaeo- and/or archicortex, or that it is the newest pallial organization to have evolved. In the absence of a universally accepted alternative to neocortex, for the remainder of this article we use the term ‘six-layered cortex’. Although some regions of this cortical domain have fewer than six layers, we justify our use by analogy to the term tetrapods, which refers to all taxa derived from ancestral four-footed vertebrates, including snakes and whales.

Overall, the evidence indicates that there are pallial, striatal and pallidal domains in most or all vertebrates²⁴. Therefore, it is reasonable to propose that the telencephala of early fishes possessed all three domains, which were then inherited as a package by later vertebrates, and independently modified in

them. The conserved organization of striatal and pallidal domains indicates that there might be constraints on how the basal ganglia can be organized. The diverse organizations of the pallial domains indicate that there are fewer constraints on how the pallium can be organized. This view has important implications for our understanding of neural mechanisms of cognition.

Avian cognition and brain function

On the basis of this new understanding of avian brain organization and its evolutionary relationships, we estimate that, as in mammals, the adult avian pallium comprises about 75% of the telencephalic volume (FIG. 1c; calculated from sagittal series of pigeon and zebra finch brain sections). This realization of a relatively large and well developed avian pallium that processes information in a similar manner to mammalian sensory and motor cortices sets the stage for a re-evaluation of the cognitive abilities of birds, which, since the 1950s, have been increasingly appreciated as far more complex than originally presumed^{91,92}. For example, pigeons can memorize up to 725 different visual patterns⁹³, learn to categorize objects as ‘human-made’ versus ‘natural’⁹⁴, discriminate cubistic and impressionistic styles of painting⁹⁵, communicate using visual symbols⁹⁶, rank patterns using transitive inferential logic⁹⁷ and occasionally ‘lie’^{98,99}. New Caledonian crows make tools out of leaves or novel human-made material, use them appropriately to retrieve food and are thought to pass this knowledge on to other crows through social learning^{100,101}. Magpies develop an understanding of object constancy at an earlier relative age in their lifespan than any other organism tested and can use this skill to the same extent as humans¹⁰². Scrub-jays show episodic memory — the ability to recall events that take place at a specific time or place, which was once thought to be unique to humans¹⁰³. This same species modifies its food-storing strategy according to the possibility of future stealing by other birds and, therefore, exhibits a behaviour that would qualify as theory-of-mind¹⁰⁴. Owls have a highly sophisticated capacity for sound localization, used for nocturnal hunting, that rivals that of humans and that is developed through learning⁶⁶. Parrots, hummingbirds and oscine songbirds possess the rare skill of vocal learning¹⁰⁵. This trait is a prerequisite in humans for spoken language and, with the exceptions of cetaceans and possibly bats, is not found in any other mammal¹⁰⁶. In addition, parrots can learn human words and use them to communicate reciprocally with humans. African grey parrots, in particular, can use human words in numerical

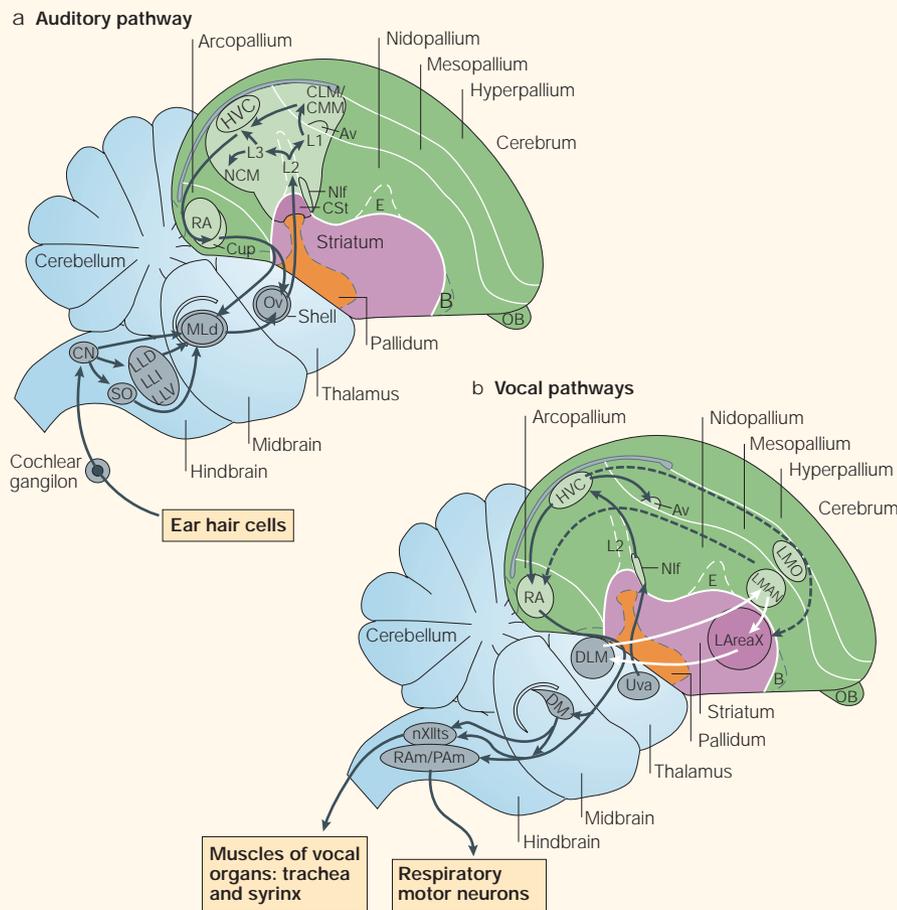


Figure 3 | Auditory and vocal pathways of the songbird brain within the context of the new consensus view of avian brain organization. Only the most prominent and/or most studied connections are indicated. **a** | The auditory pathway. Most of the hindbrain connectivity is extrapolated from non-songbird species. For clarity, reciprocal connections in the pallial auditory areas are not indicated. **b** | The vocal pathways. Black arrows show connections of the posterior vocal pathway (or vocal motor pathway), white arrows indicate the anterior vocal pathway (or pallial–basal ganglia–thalamic–pallial loop) and dashed lines show connections between the two pathways. Av, avalanche; B, basorostralis; CLM, caudal lateral mesopallium; CMM, caudal medial mesopallium; CN, cochlear nucleus; CSt, caudal striatum; DLM, dorsal lateral nucleus of the medial thalamus; DM, dorsal medial nucleus; E, entopallium; HVC (a letter-based name); LAreaX, lateral AreaX of the striatum; LLD, lateral lemniscus, dorsal nucleus; LLI, lateral lemniscus, intermediate nucleus; LLV, lateral lemniscus, ventral nucleus; LMAN, lateral magnocellular nucleus of the anterior nidopallium; LMO, lateral oval nucleus of the mesopallium; MLd, dorsal lateral nucleus of the mesencephalon; NCM, caudal medial nidopallium; Nif, interfacial nucleus of the nidopallium; nXIIIts, nucleus XII, tracheosyringeal part; OB, olfactory bulb; Ov, ovoidalis; PAm, para-ambiguous; RA, robust nucleus of the arcopallium; RAm, retroambiguous; SO, superior olive; Uva, nucleus uvaeformis.

and relational concepts^{107,108}, abilities that were once thought to be unique to humans.

So, many birds have cognitive proficiencies that are quite sophisticated, and some birds and mammals have cognitive proficiencies that clearly exceed all other birds or mammals. As these cognitive functions are carried out by the six-layered cortex in mammals¹⁰⁹ but by nuclear pallial areas in birds^{58,66,110,111}, it is clear that the mammalian six-layered cortical architecture is not the only neuroarchitectural solution for the generation of complex cognitive behaviours. It is also clear that pallial–cortical folding is not required. Birds apparently

cannot use cortical folding because of the nuclear organization of their telencephalon; among mammals such folding is related more to absolute brain size than to behavioural complexity^{112,113}. The presence of specific brain subdivisions and connections are more important factors for the generation of behavioural complexity.

An example of how avian pallial and sub-pallial areas can interact to produce complex behaviour in the context of the new view of avian brain organization can be seen in the brain pathways that control learned vocal communication (FIG. 3). Most of the telencephalic

auditory processing areas are in the pallium, adjacent to a smaller auditory area in the striatum (FIG. 3a). Likewise, most of the telencephalic vocal control nuclei are in the pallium, with one vocal nucleus in the striatum (FIG. 3b). The vocal nuclei that are involved in the production of learned vocalizations, including human speech in parrots¹¹¹, make up a pathway that directly innervates brain-stem motor neurons (FIG. 3b, black arrows). This vocal motor pathway is similar to mammalian motor corticobulbar pathways¹⁰⁶. The vocal nuclei that are involved in the imitation of vocalizations form a pallial–basal ganglia–thalamic–pallial loop (FIG. 3b, white arrows). This vocal learning pathway is similar to mammalian cortical–basal ganglia–thalamic–cortical loops^{27,106,114}, which are involved in motor learning, sensorimotor integration and additive behaviours. Other avian sensory and motor systems that are used for cognitive behaviours share a common circuit organization with the auditory and vocal pathways^{63,64}.

Conclusion

The inaccurate evolution-based terminology for the vertebrate brain that was used throughout the twentieth century became a severe impediment to the communication of scientific discoveries and the generation of new insights. Many of the tenets on which this old view of vertebrate telencephalic evolution was based have been refuted. The problems created by this view and its associated nomenclature have now been rectified for the avian brain with a new terminology that reflects the current understanding of vertebrate brain organization, homologies, evolution and function. This new understanding should facilitate a better assimilation of scientific insights into brain function through the study of birds.

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Competing interests statement

The authors declare no competing financial interests.

Online links

DATABASES

The following terms in this article are linked online to Entrez Gene:
<http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=gene>
BDNF | *DLX1* | *DLX2* | *EAG2* | *EMX1* | *ENK* | *ERB1* | *GluR1* | *mGluR2* | *NKX2.1* | *PAX6* | *ROR-β* | *SP* | *TBR1* | *ZENK*

FURTHER INFORMATION

Avian Brain Nomenclature Exchange web site:

<http://avianbrain.org/>

Digital Anatomist Project:

<http://www9.biostr.washington.edu/da.html>

Access to this interactive links box is free online.

improving resolution have not escaped the attention of the neuroscience and neuroethics communities, the media or the broader public^{2–6}. However, are the boundaries of what this technology can and cannot achieve being effectively communicated to the public? Are its limitations understood? Are the applications of the technology viewed as useful and meaningful? Are some studies more conducive to misinterpretation than others? What are the associated risks to society? From a scientific perspective, important methodological and technical assumptions guide fMRI research. However, from the public's point of view, once research results are publicized, especially when they concern personality, self-identity and other social constructs, they are bound to interact with lay conceptions of these phenomena.

To understand this complex interaction between neuroscience and society, we focused on the coverage of fMRI — as one model of frontier neurotechnology — in the print press. We investigated how both neuroscience and the media shape the social understanding of fundamental aspects of our reality and how this, in turn, points to issues of scientific communication and public involvement in neuroscience. To this end, we frame our arguments according to three trends that we have observed in press coverage of fMRI — 'neuro-realism', 'neuro-essentialism' and 'neuro-policy' — and explore how neuroethics can attend to the related ethical, legal and social issues by promoting multidirectional communication in neuroscience.

fMRI in the public eye

The increasing investigation of cognitive and social phenomena using fMRI¹ represents a relatively new venture for neuroscience. Neuroscientists who pursue such research hope for new insights into behaviour, culture and personality. However, they face new challenges in trying to convey this knowledge meaningfully to the public. Journalists, from their purview, must report these results in an adapted communication style that differs from scientific communication and adheres to a separate set of standards⁷. This creates a context in which the wider significance of research results and efforts for public outreach intermingle with the reporting of neuroscientific findings.

To understand this context specifically in relation to neuroimaging, and to launch a discussion of these issues, we carried out a press content analysis⁸ of samples of print media coverage of fMRI. Using this method, we were able to capture salient messages about the research as they are conveyed to readers.

SCIENCE AND SOCIETY

fMRI in the public eye

Eric Racine, Ofek Bar-Ilan and Judy Illes

Abstract | The wide dissemination and expanding applications of functional MRI have not escaped the attention of the media or discussion in the wider public arena. From the bench to the bedside, this technology has introduced substantial ethical challenges. Are the boundaries of what it can and cannot achieve being communicated to the public? Are its limitations understood? And given the complexities that are inherent to neuroscience, are current avenues for communication adequate?

Functional neuroimaging techniques, such as functional MRI (fMRI) and positron emission tomography (PET), have evolved as key research approaches to studying both disease processes and the basic physiology of cognitive

phenomena in contemporary neuroscience. In the clinical domain, they carry hope for guiding neurosurgical mapping, monitoring drug development and providing new approaches to disease diagnosis and management at early, possibly even presymptomatic stages. However, issues relating to these capabilities, such as technical readiness and the possibility of disease screening in advance of effective therapeutic intervention, raise substantial ethical challenges for investigators, health care providers and patients alike. In basic neuroscience, increasing numbers of non-health-related fMRI studies that touch on our personal values and beliefs have also forced us to expand our ethical perspectives¹. The wide dissemination of this research, growing applications of the technology and continuously

Links

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DLX1

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DLX2

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EAG2

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EMX1

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GluR1

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mGluR2

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Biography

The Avian Brain Nomenclature Consortium began in 1997 as a small group of neurobiologists with expertise in vertebrate brain evolution that met for dinners at the annual Society for Neuroscience meeting. Their goal was to change the inaccuracies in extant nomenclatures that had been used for almost 100 years to describe the brains of birds and other vertebrates, and which cause communication problems in the neuroscience community today. The Consortium grew to include a core group called the ThinkTankers, within a larger group called the Avian Brain Nomenclature Forum that met at Duke University, Durham, North Carolina in 2002. The members of the Forum are the authors of this report. In addition, membership of related e-mail discussion lists includes almost the entire avian neurobiology community as well as neurobiologists specializing in the study of other vertebrate brains, located in 16 different countries. This global participation ensured that current and future generations of scientists, throughout the world's universities, were a part of the effort and would rapidly institute the new nomenclature. The ThinkTank group included the following scientists (their relevant expertise is given in parentheses): Ann Butler (reptilian and fish neuroanatomy); Laura Bruce (reptilian, amphibian and mammalian neuroanatomy); András Csillag (avian neurobiology and Latin, helpful for generating a standard Latin version of the new nomenclature); Onur Güntürkün (avian visual pathways and German, helpful for translating original German brain nomenclature papers); Erich Jarvis (songbird neurobiology and molecular biology; co-organizer of the Avian Brain Nomenclature Forum); Harvey Karten (comparative avian and mammalian neuroanatomy, co-author of the pigeon brain atlas); Wayne Kuenzel (avian neuroanatomy, co-author of the chicken brain atlas); Loreta Medina (avian, reptilian and mammalian neuroanatomy); George Paxinos (mammalian neuroanatomy, co-author of widely used rat, mouse, monkey and human brain atlases); David Perkel (songbird neurobiology, comparative avian and mammalian basal ganglia function); Anton Reiner (fish, avian, reptilian and mammalian neuroanatomy, with particular expertise on the basal ganglia; co-organizer of the Avian Brain Nomenclature Forum); Toru Shimizu (avian visual pathways and brain evolution); Georg Striedter (avian and fish neurobiology, history of brain nomenclature); and Martin Wild (comparative avian and mammalian neuroanatomy). Other Consortium authors included an undergraduate student (Siang), graduate students (Dugas-Ford, Husband, Yamamoto and Yu), postdoctoral researchers (Hough, Kubikova, Smulders and Wada), and other Professors (Ball, Lee, Durand, Mello, Powers and White).

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