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Design-based stereological study of the guinea-pig (Cavia porcellus) cerebellum

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Abstract

 Guinea pigs have proved useful as experimental animal models in studying cerebellar anatomical and structural alterations in human neurological disease; however, they are also currently acquiring increasing veterinary interest as companion animals. The morphometric features of the normal cerebellum in guinea pigs have not been previously investigated using stereology. The objective of the present work was to establish normal volumetric and quantitative stereological parameters for cerebellar tissues in guinea pigs, by means of unbiased design-based stereology. Cerebellar total volume, grey and white matter volume fractions, molecular and granular layers volume fractions, cerebellar surface area, Purkinje cellular and nuclear volumes, and the Purkinje cell total count were stereologically estimated. For this purpose, cerebellar hemispheres from six adult male guinea pigs were employed. Isotropic, uniform random sections were obtained by applying the orientator method, and subsequently processed for light microscopy. The cerebellar total volume, the white and grey matter volume fractions, and the molecular and granular layer volumes were estimated using the Cavalieri's principle and the point counting system. The cerebellar surface area was estimated through the use of test lines; Purkinje cellular and nuclear volumes were analysed using the nucleator technique, whereas the Purkinje cell total count was obtained by means of

 Keywords: Guinea pig, cerebellum, stereology, neuroanatomy, nervous system.

Introduction

 The involvement of the cerebellum in motor coordination, balance and motor learning has been long and widely recognized (Brooks, 1984; Llinás and Welsh, 1993; Baillieux *et al.*, 2008; Lee *et al.*, 2015); however, a growing body of evidence involving neuroanatomical, neuroimaging and clinical studies indicates that it plays a significant role in non-motor behavioral-affective and cognitive functions, as well (Schmahmann and Caplan, 2006; Booth *et al.*, 2007; Molinari *et al.*, 2008; Cantalupo and Hopkins, 2010; Koziol *et al.*, 2011; De Smet *et al.*, 2013; Roostaei *et al.*, 2014).

 Design-based stereological techniques allow to efficiently acquire accurate and precise quantitative estimates of three-dimensional morphometric features of whole organs from measurements made on two- dimensional sections, by making use of statistical sampling and stochastic geometry principles (Boyce *et al.*, 2010).

 Most stereological investigations on the cerebellum involving laboratory animals have been carried out on mice (Woodruff-Pak, 2006; Woodruff-Pak *et al.*, 2010; Wittmann and McLennan, 2011; Kennard *et al.*, 2013; Song *et al.*, 2014), rats (Korbo *et al.*, 1993; Larsen *et al.*, 1993, 2000; Ragbetli *et al.*, 2007; Sonmez *et al.*, 2010) and rabbits (Akosman *et al*., 2011; Selçuk and Tıpırdamaz, 2020), but also on domestic animals such as

 cats (Sadeghinezhad *et al.*, 2020), pigs (Jelsing *et al*., 2006) and chicks (Tunç *et al.*, 2006). Apart from a stereological study performed on prenatal and neonatal guinea-pig cerebella following experimentally-induced intrauterine growth restriction (Mallard *et al.*, 2000), the morphometric features of the normal cerebellum in adult animals of this species have not been previously investigated using stereological techniques.

 Guinea pigs (*Cavia porcellus*) have proved useful as experimental animal models in studying cerebellar anatomical and structural alterations in human neurological disease (Lev-Ram *et al.*, 1993; Furuoka *et al.*, 2011; Čapo *et al.*, 2015; Bennet *et al.*, 2017; Cumberland *et al.*, 2017), partly due to their high degree of neurological maturity at birth in relation to the short gestation period (Altman and Das, 1967; Hargaden and Singer, 2012; Silva *et al.*, 2016), which is important for clinical studies in human medicine. Indeed, the brain of newborn guinea pigs is singularly mature, and postnatal cerebellar neurogenesis is minimal in this precocial species (Altman and Das, 1967). It was observed that, as early as 45 days through gestation, cerebellar layers in guinea-pig fetuses were distinct and well developed, with easily identifiable Purkinje cells, and with the white and gray matters well differentiated both macro- and microscopically (Silva *et al.*, 2016). Moreover, cellular proliferation events in the cerebellum, unlike other rodents, are complete at birth in the guinea pig (Lossi *et al.*, 1997).

 Recently, however, increasing interest has been addressed toward the clinical features, pathological changes and therapeutic resolution of neurological disorders of guinea pigs held as pet animals (Hollamby, 2009; Hawkins and Bishop, 2012). Most incidences of naturally-occurring cerebellar pathology reported in the literature for pet guinea pigs have an infectious etiology. Reported aetiological agents are, for instance, the lymphocitic choriomeningitis virus, leading to cerebellar hypoplasia with acute destruction of cortex folia and necrosis of granule and Purkinje cells (Monjan *et al.*, 1971; Hawkins and Bishop, 2012); *Toxoplasma gondii,* inducing granulomatous meningoencephalitis, foci of necrosis, and chronic cysts in the central nervous system (Brabb *et al.*, 2012; Gentz and Carpenter, 2012); and *Bayilisascaris procionis* larvae, causing progressive multifocal encephalomalacia and eosinophilic granulomatous inflammation of the cerebellum, midbrain and brainstem (Van Andel *et al.*, 1995).

 In light of the above-listed scientific evidence, the objective of the present work was to establish normal volumetric and quantitative stereological parameters for cerebellar tissues in adult guinea pigs, by means of unbiased design-based stereology (Gundersen and Jensen, 1987; West, 1993; Boyce *et al.*, 2010). Specifically, the present study was designed to estimate cerebellar total volume, grey and white matter volume fractions, molecular and granular layers volume fractions [by using the Cavalieri 's principle (Gundersen and Jensen, 1987)], estimate the cerebellar surface area (Schmitz and Hof, 2005), the total number of Purkinje cells [by employing the optical disector method (Gundersen, 1977; Sterio, 1984)], and the mean Purkinje cellular and nuclear volumes [through the use of the nucleator method (Gundersen *et al.*, 1988b)] in the guinea pig.

 The morphometric data emerging from the present study provide an accurate set of reference data potentially valuable as basic anatomical contribution to the field of veterinary neurology in order to help implementing the development of the diagnosis and treatment of nervous diseases in the guinea pig.

Methods

Animals and tissue preparation

140 Six adult male pet guinea pigs, weighing 569 ± 64.9 g, which spontaneously died of diseases other than those affecting the nervous system, were used for our research purposes following owners' permission. The animals did not present a history of neurological disease nor displayed pathological alterations of nervous tissues.

 According to Directive 2010/63/EU of the European Parliament and of the 22 September 2010 Council on the protection of animals used for scientific purposes, the Italian legislation (D. Lgs. n. 26/2014) does not require any approval by competent authorities or ethical committees, as this research did not influence any therapeutic decisions.

 Guinea-pig cerebella were excised from the neurocranium in their entirety, each was divided into two halves, and then immersed in a 4% phosphate-buffered formaldehyde solution to enable tissue fixation. One hemisphere of each cerebellum was randomly chosen and weighed on a digital laboratory scale. The cerebellar hemispheres were routinely processed for light microscopic examination and subsequently embedded in paraffin.

Tissue sampling and stereology

 The orientator method (Mattfeldt *et al*., 1990; Nyengaard, 1999) was applied to obtain isotropic, uniform, random sections. In essence, each cerebellar hemisphere was embedded in a paraffin block, which was placed at the center of a circle with 90 equidistant divisions along the perimeter. A random number between 0 and 90 was looked up and the paraffin medium was cut along a line parallel to the direction of the selected number. The block was placed on its cut surface at the center of a second circle, with 96 nonequidistant divisions along its perimeter. The paraffin was cut along a line parallel to the direction of a random number ranging from 0 to 96, and the block was finally re-embedded in paraffin while placed on its cut surface (Fig. 1). Consecutive 25 micrometer-thick sections were cut with a microtome at uniform constant intervals with a random start and until exhausting the organ. Every 25th section was collected using the principle of systematic uniform random sampling (Gundersen and Jensen, 1987), in order to acquire 12 to 15 sections per animal. Sections were then stained with Cresyl violet 0.1 % stain solution. A slide scanner (Optic lab H850, Plustek) was employed for capturing images from sections in order to enable the subsequent estimation of volumes and surface areas. A microscope (CX40, Olympus, Germany) equipped with an oil immersion 177 objective $(\times 100)$, connected to a microcator (MT12, Heidenhain, Traunreut, Germany) and a digital camera (MB-225) was utilized for the estimation of Purkinje cells total cellular and nuclear volumes. Geometrical probes, necessary for the stereological analysis of each structural feature represented in each section (West, 1993), were produced using a dedicated software (ImageJ; https://imagej.nih.gov).

Estimation of total and fractional volumes

 The accurate estimation of cerebellar total volume was made possible by employing cerebellar weight and transforming it into a volume, and by applying the Cavalieri's estimator, taking therefore into account tissue shrinkage. Cerebellar shrinkage secondary to histological processing allows to obtain unbiased stereological estimations insensitive to

- The estimation of total volume starting from the weight of the
- cerebellum, was performed using the following formula:
- 192 V (cerebellum) = W (cerebellum) / ρ ,
- where ρ refers to the weight-to-volume ratio of cerebellar tissue.

The estimation of the total volume of the cerebellum through use of the

Cavalieri principle was conducted by using the test point system (Fig. 2)

and following the equation below (Howard and Reed, 1998):

197
$$
V = \Sigma P \cdot SSF \cdot T \cdot (a/p)
$$

198 where ΣP is the total number of points hitting the structure; SSF (1/25) 199 represents the section sampling fraction; $T(25 \mu m)$ is the section thickness 200 and a/p (465,267 μ m²) refers to the area per point.

- 201 The fractional volume (V_v) of cerebellar structures including white matter, grey matter, molecular and granular layers, was estimated using the following formula (Gundersen *et al.*, 1988a):
- 204 V_v (structure)= $\sum P$ (structure) / $\sum P$ (cerebellum)
- 205 where $\sum P$ (structure) is the number of points hitting the white matter, gray 206 matter, molecular and granular layers, and $\sum P$ (cerebellum) is the number of points hitting the cerebellum.

 Lastly, in order to estimate the volume accounted for by each structure, each volume fraction was multiplied by the total volume of the cerebellum.

Estimation of surface area

212 The surface density (S_v) of the cerebellum was estimated by using test lines (Fig. 2b), and by employing the following formula (Howard and Reed, 1998):

215 $S_v = 2 \cdot \sum l / (\sum P \cdot l/p)$

216 Where Σ represents the total number of intersections between the outer

217 surface of the cerebellum and the test lines, ΣP refers to the points hitting

218 the molecular layer, $1/p$ (658 µm) was the length of each test line associated

to each point of the test grid.

Consequently, for estimating the surface area, surface density was multiped

by the volume of the molecular layer.

In addition, the thickness (T) of the molecular and granular layers was

calculated using the following formula (Andersen *et al.*, 2012):

224 T (layer) = V (layer) / S (layer)

where V is the volume and S is the surface area of each layer.

Estimation of Purkinje cell total count

 The optical disector method was employed for the estimation of the Purkinje cell total number, and a motorized stage designed by Department of Anatomy, Faculty of Veterinary Medicine, of the University of Tehran, Tehran, Iran, was employed for the purpose. The microscopic fields were

 selected by moving the microscope stage in the x and y directions for a constant distance spanning the entire section thickness. The unbiased counting frame principle was applied for counting the cells. The Purkinje cells whose nucleolus was located inside the counting frame or crossed the accepted lines were sampled, and those whose nucleolus came into focus within disector height were counted (Fig. 3).

The numerical density of Purkinje cells was calculated using the following

formula (Kristiansen and Nyengaard, 2012):

239 N_v (Purkinje cells) = $[\Sigma Q^{-}/(a/f \times \Sigma P \times h)] \times t/BA$

240 where ΣQ represents the total count of Purkinje cells, $a/f(9895 \mu m^2)$ is the 241 area per frame, ΣP is the total number of frames, *h* (10 µm) is the disector height, *t* is the sections mean thickness (18.5 µm), measured for each microscopic field, and BA (25 µm) is the block advance.

 Finally, for the estimation of the total number of Purkinje cells, the numerical density was multiplied by the total volume of the cerebellum, estimated using the Cavalieri's principle.

Estimation of mean Purkinje cellular and nuclear volumes

 To estimate the volumes of Purkinje cells and Purkinje cell nuclei, the nucleator technique was utilized (Gundersen *et al.*, 1988b). The volume of the sampled cells was measured by using the unbiased counting frame, and following the formula (Gundersen *et al.*, 1988b):

 $V_n = 4\pi/3 \cdot l_n^3$

253 Where l_n refers to the intercept length from the nucleolus to the border of the cytoplasm (for cellular volume), or to the border of the nucleus (for nuclear volume) of Purkinje cells.

Estimation of the coefficient of error (CE)

 The precision of the volume estimates, expressed in terms of CE, is related to the variability associated with systematic uniform random sampling (SURS) sampling and point counting of the estimator. The CE for the estimate of the volume (Gundersen and Jensen, 1987), surface area (Kroustrup and Gundersen, 1983) and Purkinje cell count [\(Braendgaard](https://pubmed.ncbi.nlm.nih.gov/?term=Braendgaard+H&cauthor_id=2332884) *et al*., 1990) was calculated.

Statistical analysis

265 All data are expressed as mean \pm standard deviation (SD). As for right-skewed distributions, a logarithmic scale was used for individual estimates of Purkinje cellular and nuclear volumes (Weber *et al.*, 1997).

Results

 All cerebella evaluated appeared normal both macroscopically and on histological examination, with all the microscopical structures being

 The mean total volume of a guinea-pig cerebellar hemisphere estimated in the present study is consistent with that calculated in a previous work, which investigated the brain morphology of domestic guinea pigs through quantitative cytoarchitectonic measurements (Kruska, 2014). Cerebellar total volume has been previously assessed by stereological techniques in other species such as humans, which exhibited a difference between sexes, with male cerebella measuring 120.5 ± 11.1 cm³ 316 in volume, while females 105.9 ± 11.2 cm³ (Taman *et al.*, 2020). Cerebellar volume has also been stereologically estimated in rabbits (Karabekir *et al.*, 318 2014) and rats (Noorafshan *et al.*, 2018), presenting volumes of 0.69 ± 0.03 319 cm³, and 0.080 ± 0.004 cm³ for each cerebellar hemisphere, respectively, but also in cats (Sadeghinezhad *et al.*, 2020), presenting a mean cerebellar 321 hemisphere volume of 2.06 ± 0.29 cm³. When comparing total cerebellar 322 volume (in cm³) in relation to body weight (in kg) in each species, it appears that the guinea pigs of the present study have a cerebellar volume to body weight ratio of 0.9, which is consistent with the 0.8 calculated for the rat (Noorafshan *et al.*, 2018), but greater than the 0.4 estimated for the rabbit (Karabekir *et al.*, 2014), and less than an approximate 1.7 for an adult individual of average weight (Taman *et al.*, 2020) and than the approximate 1.1 calculated for a medium-sized cat (Sadeghinezhad *et al.*, 2020).

 The cerebellar weight to body weight ratio was 0.1 in the guinea pig study population, which is in line with an approximate 0.13 calculated for a medium-sized cat (Sadeghinezhad *et al.*, 2020). Cerebellar volumetric modifications have been correlated with physiological factors such as age, gender (Raz *et al.*, 1998), cognitive capability, but also with several pathological neurological conditions such as Alzheimer's disease, schizophrenia and epilepsy in humans (Bottmer *et al.*, 2005; Sato *et al.*, 2007; Bas *et al.*, 2009; Andersen *et al.*, 2012). A study carried out on rats has also identified a correlation between maternal diabetes and a reduction of total cerebellar volume and thickness of all layers in the offspring (Hami *et al.*, 2016). Volumetric prediction of the cerebellum can therefore find a valuable use in further research on veterinary neurological disease affecting cognition.

 The cerebellar gray and white matter volumes have also been sterologically estimated in other species. The total gray matter volume of 345 human cerebella has been calculated to be 88.5 cm^3 , while that of the white 346 matter 22.5 cm³ (Andersen *et al.*, 2012). Cerebellar grey and white matter 347 volumes were estimated to be 1.46 ± 0.24 cm³ and 0.60 ± 0.06 cm³, respectively, for the cat (Sadeghinezhad *et al.*, 2020). When compared to the guinea pig and cat, the proportionally more voluminous grey matter in humans can be likely ascribed to their increased development of motor control, coordination, as well as cognitive functions. Moreover, it was noted that, in the early domesticated mammals such as the guinea pig, a decrease in total brain size, which is proportional to the level of encephalization of the species, along with a decrease in total cortex and areas responsible for processing sensory information and motor control,

 such as the grey matter, occurred as a consequence of the domestication process, with, however, the cognitive functions not being affected by this change (Kruska, 2005; Kaiser *et al.*, 2015; Welniak-Kaminska *et al.*, 2019). The volumes of the grey and white matter calculated in the present study are markedly greater than those reported by Mallard *et al.* (2000) for neonatal guinea pigs, which is likely due to the large age and body weight discrepancy. The influence of the physiological process of aging on volumetric changes in the cerebellar gray and especially the white matter has been assessed in several studies (Jernigan *et al.*, 2001; Walhovd *et al.*, 2005).

 Several human neurological diseases affecting cognition have also been observed to cause volume losses of the cerebellar gray and white matters (Fennema-Notestine *et al.*, 2004; Anderson *et al.*, 2009), as evidence of the role that the cerebellum plays in cognition.

 The mean volumes of the molecular and granular layers in the guinea pig cerebellum estimated in the present work are significantly greater than the corresponding values reported by Mallard *et al.* (2000) for neonatal guinea pigs, and, comparing the two studies, the volumes of the two layers are apparently not proportionally related to body weight. The mean 375 corresponding volumes referring to humans are 54.4 cm^3 for the molecular 376 layer, and 37.9 cm^3 for the granular layer (Andersen, 2004). The mean

 volume of the molecular and granular layers of the cerebellum of normal 378 rats was reported to be 0.035 cm^3 and 0.024 cm^3 , respectively (Dortaj *et al.*, [2018\)](https://www.ncbi.nlm.nih.gov/pubmed/?term=Anvari%20M%5BAuthor%5D&cauthor=true&cauthor_uid=29942437). In cats' cerebella, the mean molecular layer volume had been 380 reported to be 0.89 ± 0.16 cm³, while that of the granular layer 0.56 ± 0.10 381 cm³ (Sadeghinezhad *et al.*, 2020). The relative proportions of the molecular and granular layers of the cerebellum in the different species seem to be conserved, thanks to the similar cerebellar microscopical anatomy. As a matter of fact, the histological examination of the guinea pig cerebella permitted the clear identification of the molecular, Purkinje and granular layers with their characteristic cellular populations. The conserved volumetric trend seems to be therefore related to function.

 A stereological study carried out on intrauterine growth-restricted guinea pigs secondary to placental insufficiency in the second half period of pregnancy, has been seen to cause a reduction in the volume of the molecular and granular layers, as well as in that of the white matter in prenatal guinea-pig cerebella, therefore causing cognitive, motor and behavioral deficits in the post-natal life (Mallard *et al.*, 2000).

 When analyzing the distribution of the thickness of the molecular and granular layers in the different subjects comprising our study population, it appears that the measurements are quite consistent and

 regular, in contrast with what Sultan and Braitenberg (1993) had reported for smaller mammalian species. Andersen (2004) calculated a mean 399 thickness of the molecular layer of 590.00 ± 0.08 µm and 410.00 ± 0.15 µm for the granular layer in human cerebella. Sadeghinezhad *et al.* (2020), on 401 the other hand, calculated 133.5 ± 10.1 µm for the molecular layer and 84.7 402 ± 17.3 µm for the granular layer in cats' cerebella. Consistently with human and cats' cerebella, the molecular layer appears thicker than the granular, although not in a statistically significant manner; however, it seems that the thickness in guinea pigs is more uniformly-distributed between the two layers when compared to cats and humans' data. This can be explained by different physiological factors such as age. Indeed, a study carried out on cats' cerebella showed that aging causes an increase in granular layer thickness at the expense of that of the molecular layer (Zhang *et al*., 2006).

 With regard to the measurement of the cerebellar surface area, the ratio of cerebellar surface area to cerebellar weight in the different animals comprising the study population remains fairly constant, supporting the proportionality between cerebellar area and cerebellar weight hypothesized by Sultan and Braitenberg for larger mammals (1993), which is probably due, unlike other smaller mammalian species, to the equally constant distribution of grey matter thickness values in our guinea pig population. Further studies on larger population samples are needed to confirm this

 finding. The average surface area of the human cerebellum has been 419 previously estimated by different authors to be 550 cm^2 (Henery and 420 Mayhew, 1989), 1027 cm² (Andersen *et al.*, 2012) and 1160 cm² (Andersen *et al*., 1992). The human cerebellum, during evolution, underwent a significant expansion of its surface area both in absolute terms as well as in relation to the neocortex; this growth played a critical role in human cognitive development in comparison with other animals, given the role of the cerebellum in cognition (Barton and Venditti, 2014). In the animal kingdom, therefore, it is likely that the cerebellar surface area of highly encephalized species such as higher primates might show a greater development in comparison with mammals of a similar size. On the other hand, a mild but significant reduction in the total cerebellar area has been described in humans with advancing age, showing varying decline trends in the different vermian lobules (Raz *et al.*, 1998). A study carried out on experimentally vitamin C-deprived guinea pig fetuses has revealed a significant reduction in cerebellar surface area due to the obliteration of fissures and the fusion of opposing folia, resulting in a macroscopically- visible cerebellar dysplasia in terms of flattening of its surface, analogously to that observed in Lyssencephaly Type 2 (Čapo *et al.*, 2015). The mentioned study is of clinical relevance in pet guinea pigs due to their natural incapability of endogenous vitamin C synthesis (Nishikimi *et al.*,

 1992), analogously to humans, resulting in the necessity of its dietary supplementation, with the risk of developing vascular as well as neurological disease in case of deprivation.

442 Purkinje cells with a perikaryon volume of $3210.1 \mu m^3$ and with a 443 nuclear volume of 470.9 μ m³ were found to have the highest frequency of occurrence in the guinea pig cerebellum. Mean Purkinje cellular perikaryon 445 volumes had been estimated to be $12400 \mu m^3$ in humans (Korbo and Andersen, 1995), 4900 μ m³ (Korbo and Andersen, 1995) and 5600 μ m³ 447 (Sørensen *et al.*, 2000) in rats, 17600 μ m³ in adult minipigs (Jelsing *et al.*, 448 2006), 2207 μ m³ in rabbits (Akosman *et al.*, 2011), and 6994 μ m³ in cats (Sadeghinezhad *et al.*, 2020). If considering a mean weight for an adult individual of each species, and calculating a ratio of Purkinje volume to body weight, these findings suggest a non-allometric correlation. Indeed, the mini-pig (Jelsing *et al.*, 2006) has a Purkinje volume to body weight ratio that is six times greater than that of humans (Korbo and Andersen, 1995), whereas rodents such as the rat (Sørensen *et al.*, 2000) and the guinea pig have, respectively, ratios that are 40 and 180 times proportionally greater than that of humans. The variability encountered might be explained by the different degrees of tissue shrinkage (Andersen *et al*., 1992), by the immersion time of the tissue in the fixative (Jelsing *et al*., 2006), by the degree of postnatal development of Purkinje perikaryon volume (Jelsing *et al*., 2006), or by different degrees of significance of Purkinje cell roles in motor, sensory and cognitive functions among the different species.

 The mean total number of Purkinje cells calculated in the present work is consistent with the value reported for the whole cerebellum in a previous work carried out on neonatal guinea pigs, that is in the order of 500,000 (Mallard *et al.*, 2000). It has been demonstrated that the brain of newborn guinea pigs, species characterized by its precocity, presents a high degree of neurological maturity, and that postnatal cerebellar neurogenesis is minimal (Altman and Das, 1967). As a matter of fact, all cerebellar layers, including Purkinje cells, as well as white and gray matters, are well developed and differentiated as early as 45 days post conception (Silva *et al.*, 2016), and that all cerebellar cell proliferation events are entirely complete at birth in this species, unlike other similar rodents (Lossi *et al.*, 1997). In the adult mini-pig cerebellum, on the other hand, the total number of Purkinje cells was in the order of 2.8 million (Jelsing *et al*., 2006). The numerosity of the Purkinje cell count in the above-mentioned study was, indeed, partially explained by a significant postnatal development in total Purkinje cell number and perikaryon volume, as it had also been demonstrated in rats (Altman and Bayer, 1978), humans (Miyata *et al*., 1999), and cats (Vastagh *et al*., 2005). The total number of Purkinje cells in the whole adult rat cerebellum was estimated at around 320,000 cells (Sonmez *et al.*, 2010), which is markedly less than the value obtained for the guinea pig, and that could be explained by the complex heterogeneity of guinea pigs' Purkinje cells. It has been noted that Purkinje cells in the guinea pig cerebellum show a complex expression pattern of zebrin II, an immunohistochemical marker of cerebellar compartmental heterogeneity, showing three levels of zebrin II expression (Larouche *et al.*, 2003), as opposed to rats, where zebrin II expression only distinguishes two classes of Purkinje cells (Brochu *et al.*, 1990).

 The hypothesis that less voluminous brains tend to have a higher cellular density than larger brains (Mwamengele *et al*., 1993) does not seem to always be applicable, as is the case with the higher count of Purkinje cells in the guinea pigs comprising the present study when compared with the values reported for the rat (Sonmez *et al.*, 2010). Reports of acquired cerebellar degenerative disease in pet guinea pigs, mostly secondary to an infectious etiology, have been described in the literature, with ataxia and loss of voluntary motor control being common clinical signs, and meningoencephalitis and cerebellar cortical hypoplasia with necrosis of granule and Purkinje cells the principal histopathological findings (Monjan *et al.*, 1971; Van Andel *et al.*, 1995; Brabb *et al.*, 2012; Gentz and Carpenter, 2012; Hawkins and Bishop, 2012).

Conflict of interests

The authors have no conflict of interests to declare.

Author contributions

M.D.S.: acquisition of data, data analysis/interpretation, drafting of the

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887 Table 1. Stereological data for total volume of cerebellar hemisphere and

888 proportional volume of gray matter and white matter in six guinea pigs.

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916 Table 2. Stereological data for surface area, volume and thickness of molecular and granular layers in cerebellar hemisphere in six guinea

molecular and granular layers in cerebellar hemisphere in six guinea pigs.

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930 Table 3. Stereological data for numeral density and total number of 931 Purkinje cells in cerebellar hemisphere in six guinea pigs.

Purkinje cells in cerebellar hemisphere in six guinea pigs.

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961 Table 4. The mean coefficient of error (CE) and coefficient of variation

 Figure 1. Isotropic, uniform random sections of the guinea-pig cerebellar hemispheres were obtained by applying the orientator method. (a): A randomly chosen cerebellar hemisphere for each animal was embedded in a paraffin block and placed at the centre of a circle with 90 equidistant divisions along the perimeter. A random number between 0 and 90 was looked up and the paraffin medium was cut along a line parallel to the direction of the selected number (here, 75). (b): The block was placed on its cut surface at the center of a second circle, with 96 nonequidistant divisions along its perimeter. The paraffin was cut along a line parallel to the direction of a random number ranging from 0 to 96 (here, 50), and the block was finally re-embedded in paraffin while placed on its cut surface, 991 and consecutive 25 μ m-thick sections were cut with a microtome.

 Figure 2. Estimation of cerebellar volume and surface area by employing the point-counting and the test-lines systems. (a): The volume of the cerebellar structures was estimated by randomly superimposing a point- counting probe onto each section. The upper right corner of each point (arrow) was taken as a reference for the count of the number of points hitting the region of interest. (b): The surface area of the cerebellum was estimated by superimposing test-lines onto each section. The arrowheads show two points hitting the molecular layer, whereas the arrows indicate the intersection between test lines with the outer cerebellar surface.

 Figure 3. Use of the optical disector technique for the Purkinje cell count and of the nucleator technique for the estimation of the Purkinje cellular and nuclear volumes. **(a):** The microscopic fields were selected by moving the microscope stage in the x and y directions for a constant distance. Then, the stage of microscope moved in z-axis and the consecutive focal planes 1009 were evaluated within optical disector height (10 μ m from -5 to -15 μ m).

 (b-f): The unbiased counting frame principle was applied for the Purkinje ell count. The cells whose nucleolus was located inside the counting frame or crossed the accepted lines were sampled, and those whose nucleolus came into focus within disector height were counted. (g): The intercept length from the nucleolus to the border of the cytoplasm, or to the border of the nucleus, was measured for the estimation of Purkinje cellular and nuclear volumes, respectively.

Figure 4. Graphs showing the frequency distribution of the Purkinje

