

## An Easier Method to Analyze Stereologically the Pig's Hippocampus

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### Abstract

Most neuro-pathophysiological research involving the hippocampus uses rodents as a nonhuman model, due to the extensive experimental descriptive literature and favorable cost. The findings from the rodent hippocampus may not be generalizable to humans, though, as ontogenetically they have different hippocampal components. While the rodent model has these limitations, the use of nonhuman primates is often not feasible due to economic and ethical considerations. Pigs are a translational alternative due to the anatomic similarity of the hippocampus in humans and pigs. **Materials and Methods:** Eight pigs' brains were harvested and then analyzed according to our previously established technique. Five brains were frozen and three were stored in formalin. All eight brains were then sent to an independent histology service, where they were sectioned according to the methodology established by Holm 1994. The slabs were 10 µm with 2.5 cm<sup>2</sup> of hippocampus cross-sectional area. **Results:** The mean total hippocampus volume was 892.84 mm<sup>3</sup> ± 198.91 mm<sup>3</sup> using Holm's methodology. The mean number of cells per sample (20X magnification settings) was 9996.75, using automated ImageJ cell counting. **Discussion:** In this study, the counts of hippocampus cells were divided into two regions of interest: CA1 and CA3. Our results show that the mean number of hippocampus cells observed was 5.75 million and 2.25 million, in the CA1 and CA3 regions respectively. Holm reported 4.12 million cells in the CA1 region and 1.51 million cells in the CA3 region. The results presented here indicate the CA1 and CA3 cell percentages being 23% and 9% respectively, which are similar to the percentages reported by Holm (21% and 12%). **Conclusion:** These results corroborate previous findings and demonstrate a novel and cost-effective way to study the hippocampus of pigs in translational neurological research.

**Keywords:** neuroscience; stereological method; hippocampus; pigs**Abbreviations:** CA-Cornus Ammonis

### Introduction

Most neuro-pathophysiological research involving the hippocampus uses rodents as a nonhuman model, due to the extensive experimental descriptive literature and favorable cost. The findings from the rodent hippocampus may not be generalizable to humans, though, as ontogenetically they have different hippocampal components. Moreover, the hippocampus proportions and architectonics are significantly different between rodents and humans. While the rodent model has these limitations, the use of nonhuman primates is often not feasible due to economic and ethical considerations. Pigs are a translational alternative due to the anatomic similarity of the hippocampus in humans and pigs. In addition, pigs are a well-accepted animal research model and have lower cost when compared to nonhuman primates.

Human-size pigs have been used in research due to their anatomical and physiological similarity to humans. [1,2] The pig's brain has a gyrencephalus similar to humans, which facilitates surgical procedures and interventions. Pigs are genetically relatively homogeneous, similar to inbred laboratory rats, leading to a minimal interindividual variation.

A publication by Holm in 1994 provided a quantitative description of fundamental structural parameters, regional volumes and neuron numbers in the hippocampus of the domestic pig [3]. The volumetric and numerical data presented by Holm provided a unique opportunity to evaluate structural differences in homologous hippocampus areas between pigs and humans, and obtain a better understanding of the functional consequences of the differences in size. Despite this strong foundation, the pig hippocampus has not been clearly established as the translational model of choice for neurological studies.

In 2016, van Dijk published a systematic review comparing the hippocampus in 18 different species, where one pig study was reported [4].

The hippocampus is well recognized as the hub of neuroplasticity and neurogenesis in the central nervous system. Many diseases begin in areas CA1 and CA3 before spreading to the cortex. As an example, Masurkar, 2018 explains in great detail the progression of Alzheimer's disease, where the CA1 area is the starting point [5]. Recently, there has been an increased interest in derangements in the CA1 and CA3 areas as the basis for important clinical syndromes such as memory impairment and spatial recognition deficit [5]. Historically, many researchers have been using rodents to analyze the hippocampus, but this poses significant translational limitations, as discussed above.

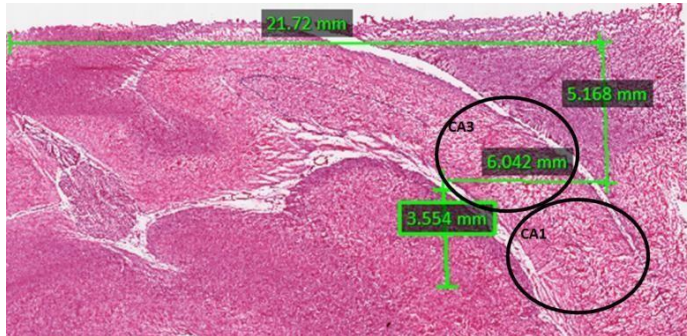
Herein, we report work that can serve as a basis for pig-based neuroscience research. The aim of this study is to extend previous work by Holm with a focus on CA1 and CA3 volume and cell count

### Materials and methods

Animal Care and Ethical approvals were obtained, and post-euthanasia surgical procedures were conducted at UBC Research Centre in Vancouver, Canada. Eight pigs' brains were harvested and then analyzed according to our previously established technique [6]. Five brains were frozen and three were stored in formalin. All eight brains were then sent to an independent histology service, where they were sectioned according to the methodology established by Holm 1994. The slabs were 10 µm with 2.5 cm<sup>2</sup> of hippocampus cross-sectional area. The hippocampus cell calculations were made according to Holm's formula:

$$\text{Hippocampus Cells (N)} = \text{Cells per slab} * 2500$$

According to the previous work by Holm, the formula above was determined using the number of cells per slab multiplied by the size of the hippocampus extension. To replicate this work, the ITCN ImageJ software plugin was used to count the number of cells per slab. Next, the ImageJ ruler was used to measure the area of the hippocampus. Holm's formula was applied to obtain the final number of hippocampal cells per subject per area. Finally, the CA1 and CA3 percentages were calculated applying the same formula on the respective areas (Figure 1).



**Figure 1:** Hippocampus measurements using ImageJ. Black ellipses represent two hippocampus areas of interest.

## Results and Discussion

The mean total hippocampus volume was  $892.84 \text{ mm}^3 \pm 198.91 \text{ mm}^3$  using Holm's methodology. The mean number of cells per sample (20X magnification settings) was 9996.75, using automated ImageJ cell counting. The percentages of hippocampal cells in areas CA1 and CA3 were 9% and 23% respectively, after normalizing by weight. Results are listed in table 1.

Hippocampus Volume	$892.84 \text{ mm}^3 \pm 198.91 \text{ mm}^3$
Mean Number of Cells per sample (20X magnification settings)	9996.75
Mean Number of Total Hippocampus cells counted per subject	24.99 million
CA1 Mean Number of cells per subject	5.75 million
CA3 Mean Number of cells per subject	2.25 million
CA1 Mean cell percentage per subject	23%
CA3 Mean cell percentage per subject	9%

**Table 1. Results**

Hippocampus Volume  $892.84 \text{ mm}^3 \pm 198.91 \text{ mm}^3$

Mean Number of Cells per sample (20X magnification settings)  
9996.75

Mean Number of Total Hippocampus cells counted per subject 24.99 million

CA1 Mean Number of cells per subject 5.75 million

CA3 Mean Number of cells per subject 2.25 million

CA1 Mean cell percentage per subject 23%

CA3 Mean cell percentage per subject 9%

The cytoarchitecture of the pig hippocampus is similar to humans. Thus, it is crucial to establish a stereological number of hippocampal cells in pigs as a basis for future studies.

It is important to correct the cell counts in proportion to body weight, as we did, and as Holm did.

Our results showed an average number of hippocampus cells of 24.99 million. This is similar to those reported by Holm (21.61 million).

In this study, the counts of hippocampus cells were divided into two regions of interest: CA1 and CA3.

Our results show that the mean number of hippocampus cells observed was 5.75 million and 2.25 million, in the CA1 and CA3 regions respectively. These results are comparable to previous studies. Holm reported 4.12 million cells in the CA1 region and 1.51 million cells in the CA3 region. The results presented here indicate the CA1 and CA3 cell percentages being 23% and 9% respectively, which are similar to the percentages reported by Holm (21% and 12%). It is important to highlight that the proportion of cells is similar in the human hippocampus.

This study provides pig hippocampus data to be used in future studies. So far, only one paper was found to address this topic. [4] However, the use of open source software to stereologically study the hippocampus was not mentioned as part of the reported methodology. The use of open source software (in our case, ImageJ) is a novel and cheaper approach to stereologically analyze the brain tissue.

Limitations: On this study we did not analyze pigs' diets, which might change the hippocampus structure and volume [7] being a variable to be considered in future comparative studies.

## Conclusion

These results corroborate previous findings and demonstrate a novel and cost-effective way to study the hippocampus of pigs in translational neurological research.

## Acknowledgements

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