Received: 27 January 2020 | Accepted: 25 May 2020

DOI: 10.1111/evj.13305

GENERAL ARTICLE

Equine Veterinary Journal WILEY

Localisation of cannabinoid and cannabinoid-related receptors in the equine dorsal root ganglia

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Funding information
This study received a grant from Formula Swiss AG, Switzerland.

Abstract

Background: Growing evidence recognises cannabinoid receptors as potential therapeutic targets for pain. Consequently, there is increasing interest in developing cannabinoid receptor agonists for treating pain. As a general rule, to better understand the actions of a drug, it would be of extreme importance to know the cellular distribution of its specific receptors. The localisation of cannabinoid receptors in the dorsal root ganglia of the horse has not yet been investigated.

Objectives: To localise the cellular distribution of canonical and putative cannabinoid receptors in the equine cervical dorsal root ganglia.

Study design: Qualitative and quantitative immunohistochemical study.

Methods: Cervical (C6-C8) dorsal root ganglia were collected from six horses (1.5 years of age) at the slaughterhouse. The tissues were fixed and processed to obtain cryosections which were used to investigate the immunoreactivity of canonical cannabinoid receptors 1 (CB1R) and 2 (CB2R), and for three putative cannabinoid-related receptors: nuclear peroxisome proliferator-activated receptor alpha (PPARα), transient receptor potential ankyrin 1 (TRPA1) and serotonin 5-HT1a receptor (5-HT1aR).

Results: The neurons showed immunoreactivity for CB1R (100%), CB2R (80% ± 13%), PPARα (100%), TRPA1 (74% ± 10%) and 5-HT1aR (84% ± 6%). The neuronal satellite glial cells showed immunoreactivity for CB2R, PPARα, TRPA1 and 5-HT1aR.

Main limitations: The low number of horses included in the study.

Conclusions: This study highlighted the expression of cannabinoid receptors in the sensory neurons and glial cells of the dorsal root ganglia. These findings could be of particular relevance for future functional studies assessing the effects of cannabinoids in horses to manage pain.

KEYWORDS
horse, CB1, CB2, PPARα, TRPA1, 5-HT1aR

1 | INTRODUCTION

The dorsal root ganglia reside within the dorsal root of the spinal nerve and are mainly constituted of a cluster of primary sensory neurons. Previous research has described the dorsal root ganglia as passive neural structures which merely "supported" the physiological communication between the peripheral nervous system and the central nervous system [1]. Nonetheless, novel evidence shows that the dorsal root ganglia neurons play a critical role in carrying sensory messages from various receptors, including those for pain and temperature, and transmitting them to the spinal cord [2].

The cell bodies of the dorsal root ganglia neurons are separated from each other by an envelope of satellite glial cells (SGCs), which play important roles in both healthy and pathological states. Since the SGCs carry receptors for numerous neuroactive agents, they have a plethora of roles including receiving signals from other cells and influencing neighbouring cells and other dorsal root ganglia neurons. These mechanisms likely influence signal processing and transmission within the dorsal root ganglia [3] and possibly contribute to the sensitisation of pain transmission nociceptors [4]. Altogether, this evidence suggests that the study of the dorsal root ganglia can have significant clinical applications for pain modulation and novel targeted therapeutic treatment [5].

A growing body of literature demonstrates that cannabinoid receptors play a critical role in nociception also through peripheral mechanisms [6]. Specifically, upon activation by medical cannabis, beneficial effects have been recorded on pain perception in humans [7-9].

Even though limited empirical research has been carried out concerning the use of medical marijuana for pain treatment in domestic animals [10] and horses [11], the use of cannabis products in animals is expanding [12]. Of the cannabis products, cannabidiol (CBD), a non-psychoactive compound found in *cannabis sativa*, seems to be one of the most promising therapeutic substances. Due to its numerous health-related benefits, CBD has found multiple clinical applications in the medical field, including analgesic, anti-inflammatory, anti-spasmodic and anti-anxiety uses [13,14]. For many years, it was assumed that the beneficial effects of the cannabinoids were mediated exclusively by cannabinoid receptors 1 (CB1R) and 2 (CB2R). However, it is currently known that phytocannabinoids may act on multiple targets outside the endocannabinoid system, such as other G-protein-coupled receptors (GPRs), the transient receptors potential (TRPs) channel, nuclear peroxisome proliferator-activated receptors (PPARs), and serotonin receptors [15]. In particular, CBD, which shows indirect interaction with CB1R and CB2R, seems to be involved in the modulation of receptors, such as the serotoninergic 5-HT1a receptor (5-HT1aR), and the transient receptors potential ankyrin 1 (TRPA1) and vanilloid 1 (TRPV1), the latter two being excitatory ion channels expressed by the sensory neurons mediating somatic and visceral pain [7].

As a general rule, to better understand the effects exerted by a drug, it is important to know the cellular distribution of its specific receptors. Currently, only a small number of studies have been published regarding the expression of cannabinoid receptors in the dorsal root ganglia of animals [16-19] and no analogous studies have yet been carried out on horses.

Thus, the current study was designed to immunohistochemically localise, two canonical cannabinoid receptors (CB1R and CB2R) and three cannabinoid-related receptors (PPARα, TRPA1 and 5-HT1aR) in the equine dorsal root ganglia.

2 | MATERIALS AND METHODS

2.1 | Animals

The cervical (C6-C8) dorsal root ganglia were collected from the left and the right halves of the carcasses of six horses (four males and two females) which were slaughtered for human food purposes. The horses (1.5 years of age) were of two breeds, three Polish and three half-breeds. The complete cell blood count (CBC) and routine serum biochemical analyses were carried out with blood samples taken at the time of exsanguination. The horses were considered healthy on the basis of normal results of the CBC count and routine serum biochemical analyses. In addition, the horses did not show lameness of either the thoracic or the pelvic limbs.

The tissues were fixed and processed to obtain cryosections, as described elsewhere [20].

Since the suppliers of the primary antibodies used in the current study stated that they were rat-specific (CB2R, TRPA1 and 5-HT1aR) or reacted with rat tissues (CB1R, PPARα), the experiments on the equine dorsal root ganglia were carried out by including rat C6-C8 dorsal root ganglia as positive controls (Figure S1). The distribution of the study receptors in the subclasses of the rat sensory neurons was outside the scope of this study and was therefore not evaluated.

2.2 | Immunofluorescence

The cryosections (14-16 μm thickness), were hydrated in phosphate-buffered saline (PBS) and processed for immunostaining. To block non-specific bindings, the sections were incubated in a solution containing 20% normal goat serum (Colorado Serum Co.), 0.5% Triton X-100 (Sigma Aldrich) and bovine serum albumin (1%) in PBS for 1 hour at room temperature (RT) (22-25°C). The cryosections were incubated overnight in a humid chamber at RT with the primary antibodies (Table 1) diluted in 1.8% NaCl in 0.01 M PBS containing 0.1% sodium azide. After washing in PBS (3 × 10 minutes), the sections were incubated for 1 hour at RT in a humid chamber with the secondary antibody [goat F(ab)2 anti-rabbit FITC; ab98430] (abcam) diluted in PBS. After washing in PBS (3 × 10 minutes), to identify the dorsal root ganglia neurons and the SGCs and to determine the proportion of neurons immunoreactive for each of the markers studied, the sections were counterstained with Blue fluorescent Nissl stain solution (NeuroTrace®, N-21479, dilution 1:200) (Molecular

TABLE 1 Primary antibodies used in the study

Primary antibody	Host	Code	Dilution	Source
CB1R	Rabbit	ab23703	1:100	abcam
CB2R	Rabbit	ab45942	1:200	abcam
PPARα	Rabbit	NB600-636	1:200	Novus Biol.
5-HT1a receptor	Rabbit	ab85615	1:100	abcam
TRPA1	Rabbit	ab58844	1:100	abcam

Primary antibodies Suppliers: abcam, Cambridge, UK. Novus Biologicals, Littleton, CO, USA.

Probes). The cryosections were then washed in PBS (3 × 10 minutes) and mounted in buffered glycerol at pH 8.6. A minimum of 100 Nissl-stained neurons were counted for each marker. Data were collected from preparations obtained from four animals ($n = 4$). The percentages of immunopositive neurons were expressed as mean ± standard deviation.

2.3 | Specificity of the primary antibodies

The choice of the primary antibodies used in the study was based on the homology of the aminoacidic sequence between the immunogen of the commercially available antisera and the horse proteins, verified by the "alignment" tool available on the Uniprot database (www.uniprot.org) and the BLAST tool of the National Center for Biotechnology information (NCBI) (www.ncbi.nlm.nih.gov).

CB1R—The immunogen used to obtain antibody ab23703 was the synthetic peptide MSVSTDTSAEAL, corresponding to carboxy-terminal amino acids 461-472 of Human cannabinoid receptor I. The homology between the full amino acid sequences of Horse (F6SIU9) and Human (P21554) CB1R was 97.88%, and the correspondence with the specific sequence of the immunogen was 100%.

CB2R—The immunogen used to obtain antibody ab45942 was the synthetic peptide conjugated to keyhole limpet haemocyanin (KLH) derived from within residues 200-300 of Rat cannabinoid receptor II. The homology between the full amino acid sequences of Horse (F7CUS7) and Rat (Q9QZN9) CB2R was 80.9%, and the correspondence with the specific sequence of the immunogen was 83.33%.

PPARα—Antibody NB600-636 was prepared from whole rabbit serum produced by repeated immunisations using a synthetic peptide corresponding to amino acids 1 to 18 of Mouse PPARα. The homology between the full aminoacidic sequences of Horse (F7DSM8) and Mouse (P23204) PPARa was 90.81%, and the correspondence with the specific sequence of the immunogen was 100%.

TRPA1—The immunogen used to obtain antibody ab58844 was peptide EKQHELIKLIIQKME corresponding to amino acids 1060-1075 of Rat TRPA1. The homology between the full amino acid sequences of Horse (F7DXW9) and Rat (Q6RI86) TRPA1 was 82%, and the correspondence with the specific sequence of the immunogen was 100%.

Serotonin 5-HT1aR—The immunogen used to obtain antibody ab85615 was the synthetic peptide corresponding to Rat 5HT1aR amino acids 100-200 conjugated to keyhole limpet hemocyanin. The homology between the full amino acid sequences of Horse (Q0EAB6) and Rat (P19327) 5HT1aR was 89.3%, and the correspondence with the specific sequence of the immunogen was 99%.

In addition, the specificity of the primary antibodies employed was also tested using Western blot (Wb) analysis.

2.4 | Specificity of the secondary antibody

The secondary antibody specificity was tested by the lack of signal after omission of the primary antibody on dorsal root ganglia tissues.

2.5 | Fluorescence microscopy

The preparations were analysed by the same observer, and the images were recorded and adjusted as described elsewhere [21].

2.6 | Western blot: specificity of the primary antibodies

Cervical (C6-C8) dorsal root ganglia and spinal cord samples were collected, frozen in liquid nitrogen, and stored at −80°C until sample processing. The primary antibodies were tested in our laboratories according to standardised protocols [21]. The Wb analysis of CB1R (1:500) revealed double bands of ~100 and 120 kDa (the theoretical molecular weight of CB1R is 53-60 kDa) (Figure 1A), that of CB2R (1:1,000) revealed a major band of ~90 kDa (the theoretical molecular weight of CB2R is 45 kDa) (Figure 1B), that of PPARα (1:2000) revealed a single band of ~50 kDa (the theoretical molecular weight of PPARα is 52 kDa) (Figure 1C), that of TRPA1 (1:500) revealed a major band at ~100 kDa (the theoretical molecular weight 100 kDa) (Figure 1D) and that of the 5-HT1aR (1:3000) revealed a major band at ~50 kDa (theoretical molecular weight 46 kDa) (Figure 1E). Western blot analysis confirmed the specificity of the anti-PPARα,-TRPA1 and-5-HT1aR primary antibodies; the latter finding is consistent with the 100% homology of the amino acid sequence between the immunogen of the antisera and the horse proteins (see above). The CB1R (100% homology) and CB2R (83% homology) showed bands with apparent double molecular weights; it is known that G-protein receptors, such as

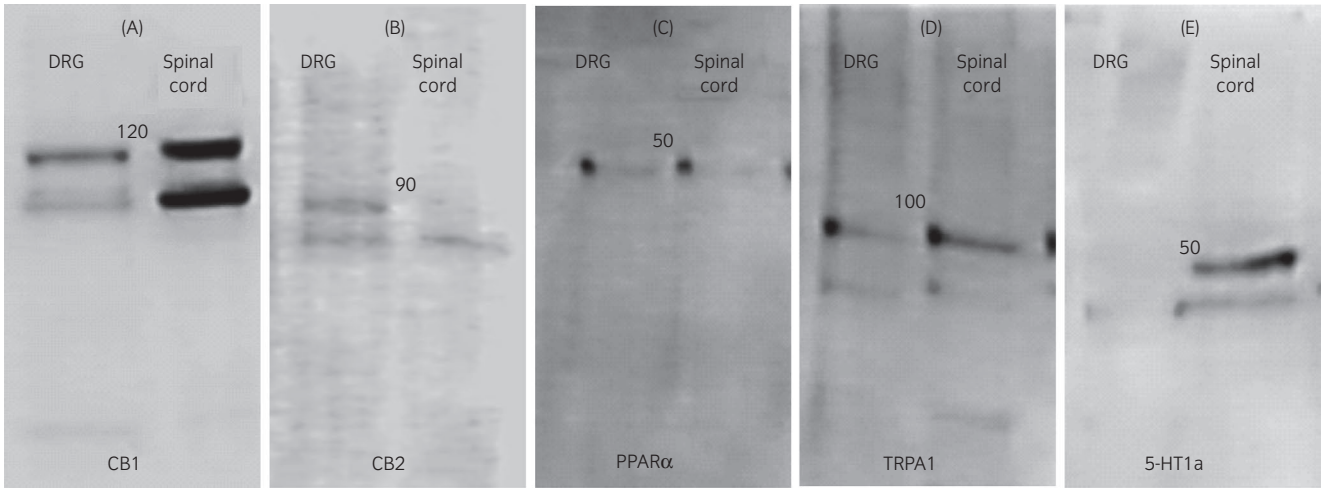

FIGURE 1 Western blot (WB) analysis showing the specificity of the following primary antibodies utilised: (A) rabbit anti-cannabinoid receptor 1 (CB1); (B) rabbit anti-cannabinoid receptor 2 (CB2); (C) rabbit anti-nuclear peroxisome proliferator-activated receptor alpha (PPARα); (D) rabbit anti-transient receptor potential ankyrin 1 (TRPA1) and e) rabbit anti-serotonin receptor 5-HT1a (5-HT1a). The antibody anti-CB1 receptor showed double bands of ~100 and 120 kDa (the theoretical molecular weight of the CB1 receptor is 53-60 kDa), more evident in the spinal cord. The antibody anti-CB2 receptor showed a double band (only in the dorsal root ganglia) of ~ 90 kDa (the theoretical molecular weight of the CB2 receptor is 45 kDa). The antibody anti-PPARα showed a single band of ~ 50 kDa (the theoretical molecular weight of PPARα is 52 kDa). The antibody anti-TRPA1 showed a major band at ~ 100 kDa (theoretical molecular weight 100 kDa). The antibody anti-5-HT1A receptor showed a double band (only in the spinal cord) at ~ 50 kDa (theoretical molecular weight 46 kDa). The numbers on the dorsal root ganglia and the spinal cord lines indicate the molecular weights. The images of the immunoblots were slightly adjusted in brightness and contrast to match their backgrounds

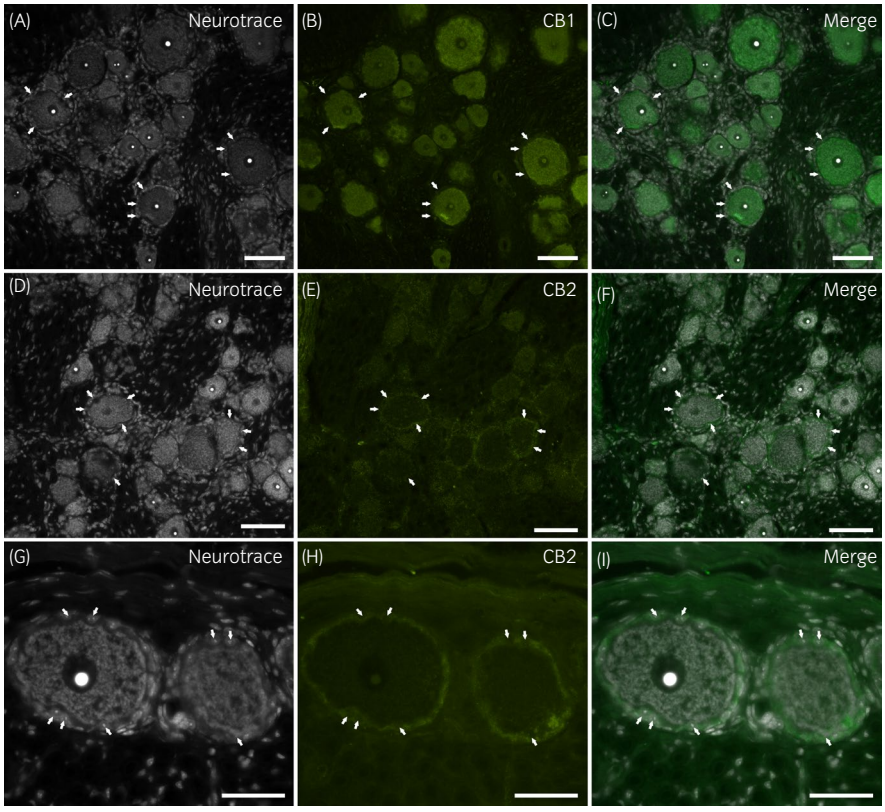

FIGURE 2 Photomicrographs of cryosections of an equine cervical (C8) dorsal root ganglion showing cannabinoid receptor 1 (CB1R) (A-C) and cannabinoid receptor 2 (CB2R) (D-I) immunoreactivity (IR). (A-C) CB1R-IR was expressed by the sensory neurons whereas the satellite glial cells, the nuclei of which are indicated by arrows, were CB1R-negative. (D-I) The arrows indicate the satellite glial cells which were CB2R immunoreactive. Sensory neurons, in particular the smallest ones, showed very faint granular CB2R immunolabelling. Scale bar = 50 μm

CB1R, may show dimerisation, and form functionally active homodimer and heterodimer complexes, and that polyclonal antibodies directed against the carboxy-terminal may only recognise the high molecular weight form of CB1R [22]. An apparent dimeric form for the other G-protein receptor CB2R was also observed; however, no studies regarding the dimeric form of this receptor

are available in the literature. Nevertheless, the possibility that the anti-CB2R antibody may recognise only the band in which the receptor is conjugated to another CB2R (or to another protein), as observed for the anti-CB1R antibody, cannot be excluded.

3 | RESULTS

3.1 | CB1R

Bright cytoplasmic and nucleolar CB1R immunoreactivity (CB1R-IR) was displayed, with different degrees of intensity, by 100% of the neurons (604/604 cells counted, *n* = 4) (Figure 2A-C); no distinction of CB1R immunolabelling was observed among neurons of different sizes. The nerve processes also showed CB1R-IR, although it was fainter than that observed in the neuronal somata. The SGCs also showed faint CB1R-immunolabelling. This finding is partially consistent with what has been observed in the rat dorsal root ganglia in which neurons and SGCs expressed faint cytoplasmic and nuclear CB1R-IR [19].

3.2 | CB2R

The CB2R-IR was expressed by 80% ± 13% of the neurons (665/799 cells counted, *n* = 4) and by the SGCs. In general, there was an inverse correlation between the brightness of CB2R-IR in the neurons and SGCs; small-to-medium-sized neurons showed brighter granular CB2R cytoplasmic immunolabelling in comparison with larger ones which were encircled by brightly labelled SGCs (Figure 2D-I). In the rat dorsal root ganglia, CB2R-IR was expressed by the neuronal nuclei [19].

3.3 | TRPA1

The TRPA1-IR was expressed by the cytoplasm (moderate staining) and nucleus (bright staining) of both the neurons (74% ± 10%; 497/698 cells counted, *n* = 4) and the SGCs, and by the nerve processes (Figure 3A-C). Of the nerve processes, thin and unmyelinated nerve fibres showed brighter TRPA1 immunolabelling than large myelinated nerve fibres. This finding is partially consistent with what

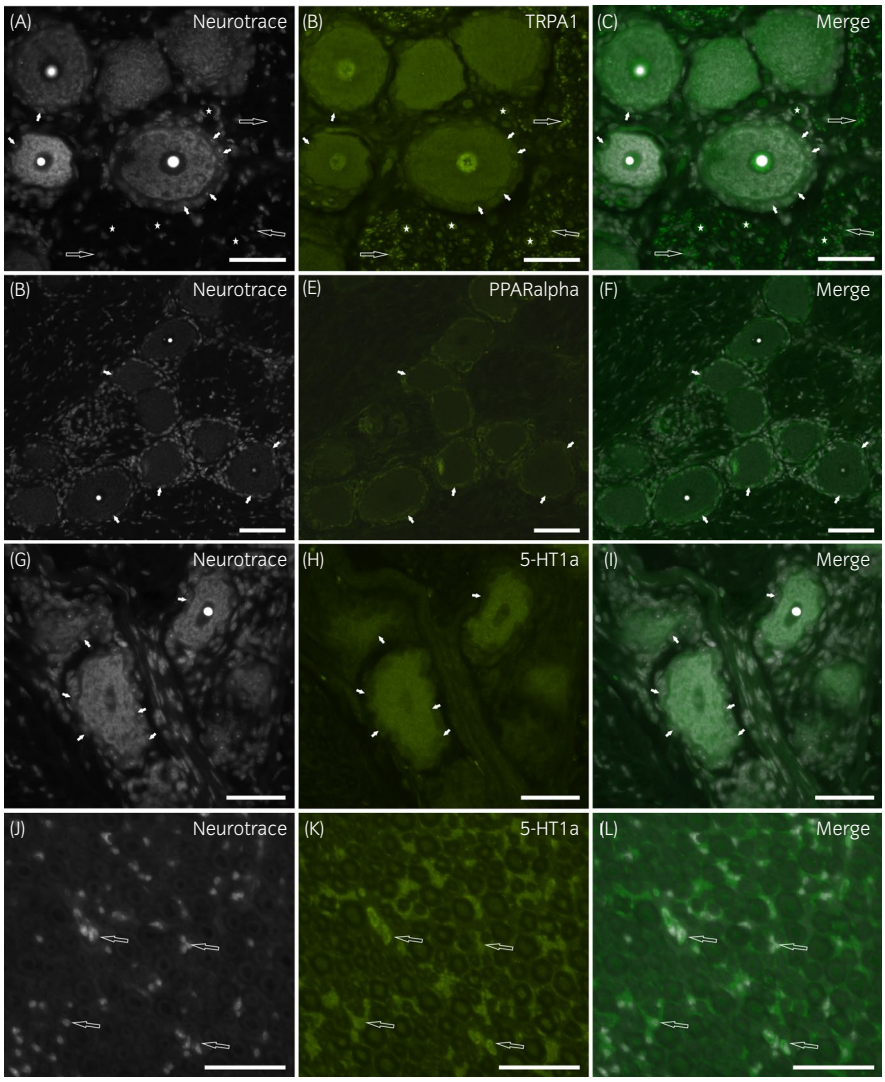

FIGURE 3 Photomicrographs of cryosections of an equine cervical (C8) dorsal root ganglion showing TRPA1- (A-C), PPARα- (D-F) and 5-HT1a receptor- (G-L) immunoreactivity (IR). (A-C) Sensory neurons and satellite glial cells (small arrows) expressed TRPA1-IR. Large arrows indicate groups of amyelinic sensory fibres which showed very bright TRPA1-IR whereas the nerve fibres with a larger diameter (stars) showed weaker immunostaining. (D-F) Weak PPARα-IR was expressed by the cytoplasm of all the neurons whereas the satellite glial cells (arrows) showed bright PPARα-IR. (G-I) Sensory neurons expressed bright 5-HT1a receptor-IR; satellite glial cells (arrows) also showed moderate 5-HT1a receptor-IR. (J-L) The Schwann cells (arrows) showed 5-HT1a receptor-IR. Scale bar = 50 μm

has been observed in the rat dorsal root ganglia in which only the neuronal cytoplasm expressed TRPA1-IR (Figure S1 A-C).

3.4 | PPARα

Faint cytoplasmic PPARα-IR was displayed by all the neurons (100%; 456/456 cells counted, *n* = 4) whereas it was brightly displayed by the cytoplasm of the SGCs (Figure 3D-F). The PPARα-IR was also expressed by the endothelial cells of the blood vessels (data not shown). This finding is partially consistent with what has been observed in the rat dorsal root ganglia in which neurons and SGCs expressed PPARα-IR [19].

3.5 | 5-HT1aR

Moderate cytoplasmic 5-HT1aR-IR was expressed by 84% ± 6% of the neurons (462/547 cells counted, *n* = 4); faint to moderate 5-HT1aR-IR was also expressed by the SGCs (Figure 3G-I) and the Schwann cells (Figure 3J-L). This finding is partially consistent with what has been observed in the rat dorsal root ganglia in which only the neuronal cytoplasm expressed 5-HT1aR-IR (Figure S1 D-F).

The results of the cellular distribution and intensity of the immunolabelling in the equine dorsal root ganglia are summarised in semiquantitative Table 2.

4 | DISCUSSION

This is the first study aimed at investigating the expression and localisation of cannabinoid receptors (CB1R and CB2R) and cannabinoid-related receptors (PPARalpha, TRPA1 and 5-HT1aR) in the equine dorsal root ganglia.

Cannabinoid CB1R is widely expressed throughout the nociceptive system and its activation by endogenous or exogenous cannabinoids modulates neurotransmitter release. Previous studies carried out on humans and rodents have shown CB1R-IR expression limited to the sensory neurons [16,19,23] whereas the present study showed that CB1R-IR is localised in both the equine sensory neurons and SGCs in line with data recently reported for dogs [18,19].

In humans, the CB2R-IR has been localised exclusively in nociceptive dorsal root ganglia neurons [23]. In the present study, CB2R-IR was observed in the equine dorsal root ganglia neurons and SGCs, as previously observed in rats [17]. The CB2R-IR has recently been observed in the neuronal nuclei of rats and dogs as well as in the Schwann cells of dogs [19]. It has been shown that the application of CB2R agonists functionally inhibits nociceptive signalling in human dorsal root ganglia neurons [24]. Based on the authors' findings, it can be hypothesised that, in horses, the same agonists could be useful for the peripheral modulation and treatment of painful sensation [24].

In addition to canonical cannabinoid receptors, multiple targets can be influenced by phytocannabinoids unravelling the complexity of this endocannabinoid system. For instance, PPARs belong to the family of intranuclear receptors which act as transcription factors, modulating different physiological functions. Once activated by their ligand, PPARs induce the expression of hundreds of genes in each cell type [25]. However, their activation has also been shown to result in rapid cellular changes which do not require transcription, including reduction of inflammation [26,27]. Recent studies have shown that cannabinoids activated PPARs [15,27,28], and that this activation is associated with some of the pain-relieving, anti-inflammatory and neuroprotective properties of cannabinoids.

In the current study, PPARα-IR expression was observed in both the sensory neurons and the SGCs as has previously been shown in rats [19] whereas PPARα was detected exclusively in the neurons or in the SGCs respectively in mice and dogs [19,29]. PPARα-IR was also observed in the endothelial cells of the dorsal root ganglia blood vessels. Recent evidence has shown that endocannabinoids exert a pro-homeostatic function on vascular biology by means of complex mechanisms involving canonical as well as putative cannabinoid receptors [30]. Specifically, different cannabinoid receptors can

TABLE 2 Semiquantitative evaluation of the density of CB1, CB2, PPARα, TRPA1 and 5-HT1a receptors immunoreactivity in different cellular elements (neurons, nerve processes, satellite glial cells, Schwann cells, blood vessels) of the equine cervical dorsal root ganglia.

	Cervical dorsal root ganglion				
Receptors	**Neurons**	**Satellite glial cells**	**Schwann cells**	**Blood vessels**	**Nerve processes**
CB1	C^D ++/+++ Nucleolar +++	N^D +	–	–	C^D+
CB2	C^D +/++	$C^{D+/}$++	–	–	–
PPARα	C^D+	C^D+++	–	E^D++	–
TRPA1	N^D+++ C^D++	N^D++	–	–	C^D++/+++
5-HT1aR	C^D++ N^D+	C^D+	C^D++	–	–

Abbreviations: C, cytoplasmic; D, diffuse labelling; E, endothelium; N, nuclear.

Immunoreactive cells were graded as: –, negative; +, faintly stained; ++, moderately stained; +++, brightly stained.

actively influence vasodilation at different cellular sites, ie nerves, endothelial cells, vascular smooth muscle cells or pericytes [31].

The TRPA1 is required for normal mechano- and chemo-sensory functions in specific subsets of vagal, splanchnic and pelvic afferents [32]. The TRPA1 also mediates somatic and visceral pain in response to stimulation of a chemical, mechanical or thermal origin [33,34], and can be desensitised by different mechanisms [35]. Data from the present study demonstrated that TRPA1 was expressed by the sensory neurons and by the SGCs, with bright immunolabelling in thin unmyelinated nerve fibres. This is partially in line with previous studies carried out on humans and rodents showing that the TRPA1-IR sensory neurons are involved in nociception [36]. As a consequence, the analgesic effects of CBD might be due, in part, to its ability to activate and desensitise the TRPA1 [37].

Another finding which emerged from this study was the expression of 5-HT1aR in sensory neurons, SGCs and Schwann cells. It is well established that serotonin (5-HT) exerts a pivotal role in sensory information processing [38]. At the level of the spinal cord, 5-HT is primarily released by the descending bulbospinal serotonergic neurons and causes analgesia by inhibiting dorsal horn neuronal responses to noxious stimuli by means of the activation of the 5-HT1aR [39]. In addition, the activation of the 5-HT1aR inhibits glutamate release from the sensory neurons, reducing pain transmission [40]. Functional studies have suggested the presence of the 5-HT1aR in the dorsal root ganglia and its role in nociception [41]. In the present study, the 5-HT1aR-IR was observed in the sensory neurons and the SGCs of horses, and in the neurons of rats. Therefore, it is plausible to consider that the 5-HT1aR might play a role in pain modulation. We also observed the 5-HT1aR in Schwann cells, a finding which is consistent with the expression of other serotoninergic receptors on Schwann cells in rodents [42]. In summary, this study showed a notable distribution of cannabinoid and cannabinoid-related receptors in the equine dorsal root ganglia.

The reduced number of animals and ganglia considered, as well as the specificity of the anti-CB2R antiserum, represents the limitations of the present study. Additional biomolecular studies as well as the neurochemical characterisation of the immunoreactive dorsal root ganglia neurons are needed to reinforce our current data. Nonetheless, these findings represent novel insight regarding the complex processing of nociceptive input in the equine peripheral nervous system and will hopefully encourage new studies regarding the role of cannabinoid-related receptors in the equine dorsal root ganglia and their interactions with non-psychotic cannabis-derived molecules. In addition, cannabinoids, specifically CBD, have been proposed as an opioid alternative in human medicine, having a comparable efficacy with a better safety profile [43].

5 | CONCLUSIONS

Cannabinoid and cannabinoid-related receptors had a wide distribution in the sensory neurons and SGCs of the equine dorsal root ganglia. These findings represented an important anatomical basis upon which it would be possible to continue with other preclinical and clinical studies aimed at investigating and possibly supporting the specific therapeutic uses of non-psychotropic cannabinoid agonists against noxius stimulation in horses.

ETHICAL ANIMAL RESEARCH

Experiments including rat tissues were approved by the National Health Authority (authorisation no. 112/2018-PR of 12 February 2018), in accordance with the DL 26/2014 and the European Union Directive 2010/63/EU, and under the supervision of the Central Veterinary Service of the University of Bologna. The equine elements of the study were performed on material obtained from an abattoir.

OWNER INFORMED CONSENT

Not applicable.

CONFLICT OF INTERESTS

No competing interests have been declared.

AUTHOR CONTRIBUTIONS

R. Chiocchetti, R. Rinnovati and F. Giancola contributed to the study design. The immunohistochemical experiments were carried out by A. Stanzani, G. Galiazzo, C. Tagliavia, M. De Silva and Y. Capodanno. R. Chiocchetti and A. Stanzani contributed to acquisition of data. All authors interpreted the data. R. Chiocchetti contributed to drafting of the manuscript. Y. Capodanno, A. Stanzani and M. De Silva contributed to revision of the manuscript. All authors contributed to the study execution and approved the final manuscript.

PEER REVIEW

The peer review history for this article is available at https://publons.com/publon/10.1111/evj.13305.

DATA AVAILABILITY STATEMENT

The data that support the findings of this study are available from the corresponding author upon reasonable request.

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SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section.

How to cite this article: Chiocchetti R, Rinnovati R, Tagliavia C, et al. Localisation of cannabinoid and cannabinoid-related receptors in the equine dorsal root ganglia. *Equine Vet J*. 2020;00:1–9. https://doi.org/10.1111/evj.13305