

The *du^{2J}* mouse model of ataxia and absence epilepsy has deficient cannabinoid CB₁ receptor-mediated signalling

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Key points

- Cerebellar ataxias are progressive debilitating diseases with no known treatment and are associated with defective motor function and, in particular, abnormalities to Purkinje cells.
- Mutant mice with deficits in Ca²⁺ channel auxiliary $\alpha 2\delta$ -2 subunits are used as models of cerebellar ataxia.
- Our data in the *du^{2J}* mouse model shows an association between the ataxic phenotype exhibited by homozygous *du^{2J}/du^{2J}* mice and increased irregularity of Purkinje cell firing.
- We show that both heterozygous *+/du^{2J}* and homozygous *du^{2J}/du^{2J}* mice completely lack the strong presynaptic modulation of neuronal firing by cannabinoid CB₁ receptors which is exhibited by litter-matched control mice.
- These results show that the *du^{2J}* ataxia model is associated with deficits in CB₁ receptor signalling in the cerebellar cortex, putatively linked with compromised Ca²⁺ channel activity due to reduced $\alpha 2\delta$ -2 subunit expression. Knowledge of such deficits may help design therapeutic agents to combat ataxias.

Abstract Cerebellar ataxias are a group of progressive, debilitating diseases often associated with abnormal Purkinje cell (PC) firing and/or degeneration. Many animal models of cerebellar ataxia display abnormalities in Ca²⁺ channel function. The ‘ducky’ *du^{2J}* mouse model of ataxia and absence epilepsy represents a clean knock-out of the auxiliary Ca²⁺ channel subunit $\alpha 2\delta$ -2, and has been associated with deficient Ca²⁺ channel function in the cerebellar cortex. Here, we investigate effects of *du^{2J}* mutation on PC layer (PCL) and granule cell layer (GCL) neuronal spiking activity and, also, inhibitory neurotransmission at interneurone–Purkinje cell (IN-PC) synapses. Increased neuronal firing irregularity was seen in the PCL and, to a less marked extent, in the GCL in *du^{2J}/du^{2J}*, but not *+/du^{2J}*, mice; these data suggest that the ataxic phenotype is associated with lack of precision of PC firing, that may also impinge on GC activity and requires expression of two *du^{2J}* alleles to manifest fully. The *du^{2J}* mutation had no clear effect on spontaneous inhibitory postsynaptic current (sIPSC) frequency at IN-PC synapses, but was associated with increased sIPSC amplitudes. *du^{2J}* mutation ablated cannabinoid CB₁ receptor (CB₁R)-mediated modulation of spontaneous neuronal spike firing and CB₁R-mediated presynaptic inhibition of synaptic transmission at IN-PC synapses in both *+/du^{2J}* and *du^{2J}/du^{2J}* mutants, effects that occurred in the absence of changes in CB₁R expression. These results demonstrate that the *du^{2J}* ataxia model is associated with deficient CB₁R signalling in the cerebellar cortex, putatively linked with compromised Ca²⁺ channel activity and the ataxic phenotype.

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Abbreviations CB₁R, cannabinoid CB₁ receptor; CV, coefficient of variation; GC, granule cell; GCL, granule cell layer; GPCR, G protein-coupled receptor; IN-PC, interneurone–Purkinje cell; ISI, inter-spike interval; PC, Purkinje cell; PCL, Purkinje cell layer; WIN55, WIN55,212–2.

Introduction

Cerebellar ataxias comprise a group of progressive diseases associated with motor incoordination and are typically associated with dysfunction and/or degeneration of PCs, which represent the sole efferent output of the cerebellar cortex. A number of mutant mouse models exhibit specific ataxias with diverse behavioural phenotypes at different developmental stages (Green, 1981; Grüsser-Cornehls & Baurle, 2001), including the *du^{2J}* mutation that exhibits behavioural traits consistent with cerebellar ataxia and absence epilepsy. *du^{2J}* mice have mutations in the *Cacna2d2* gene which encodes the $\alpha 2\delta$ -2 auxiliary Ca²⁺ channel subunit (Donato *et al.* 2006), one of four $\alpha 2\delta$ subunit isoforms ($\alpha 2\delta$ -1–4) that exert auxiliary effects on Ca²⁺ channel biophysical properties and physiological function (Gao *et al.* 2000; Hobom *et al.* 2000; Klugbauer *et al.* 2003; Bauer *et al.* 2010; Dolphin, 2012; Hoppa *et al.* 2012). *du^{2J}* mice are part of a group of mutant mouse strains together with either spontaneous (*Cacna2d2^{entla}* and *Cacna2d2^{du}* alleles) or targeted (*Cacna2d2^{tm1NClF}*) $\alpha 2\delta$ -2 disruptions, all of which typically exhibit smaller than normal size, comparable ataxia phenotypes, absence seizures and paroxysmal dyskinesia (Barclay *et al.* 2001; Brodbeck *et al.* 2002; Inanov *et al.* 2004; Brill *et al.* 2004; Donato *et al.* 2006; Walter *et al.* 2006). The *Cacna2d2^{entla}* allele predicts a full-length protein with an inserted region in the $\alpha 2$ moiety of $\alpha 2\delta$ -2 and is associated with reduced PC Ca²⁺ currents (Brill *et al.* 2004). The *Cacna2d2^{du}* allele disrupts *Cacna2d2* in intron 3, yielding a truncated $\alpha 2\delta$ -2 protein and resulting in reduced native and recombinant Ca_v2.1 Ca²⁺ channel expression (Barclay *et al.* 2001; Brodbeck *et al.* 2002). The *Cacna2d2^{du2J}* allele used here has a 2 bp deletion in exon 9 of *Cacna2d2* resulting in complete ablation of $\alpha 2\delta$ -2 expression and reduced PC Ca²⁺ currents (Donato *et al.* 2006). In *du* mutant mice, a reduction in Ca²⁺ influx, leading to compromised Ca²⁺-dependent K⁺ channel (SK) activity and irregular pacemaking, was proposed to underlie the ataxic phenotype (Walter *et al.* 2006); similarly, the *du^{2J}* mutation exhibits increased PC firing irregularity, although this could not be normalised using SK blockers (Donato *et al.* 2006).

Here, we extend previous studies to examine the effect of *du^{2J}* mutation on basal neuronal network activity and synaptic transmission and, further, on G protein-coupled receptor (GPCR)-mediated presynaptic inhibition of synaptic transmission in the cerebellum. In particular, CB₁ GPCRs are strongly expressed in the cerebellar cortex, where they modulate GABA transmission at IN-PC synapses to modulate PC total output (Ma *et al.* 2008;

Wang *et al.* 2011). We demonstrate that the *du^{2J}* phenotype exhibits deficient CB₁R signalling at the neuronal network level that reflects, at least in part, ablation of CB₁R modulation of inhibitory neurotransmission at IN-PC synapses, but which does not result from reduced CB₁R expression. These results suggest that $\alpha 2\delta$ -2 deficits in *du^{2J}* mutants affect GPCR-mediated modulation of inhibitory transmission in the cerebellar cortex, with consequential effects upon PC spike firing activity; such deficits may be associated with ataxic phenotypes and, potentially, contribute to disease.

Methods

Ethical approval

All work was subject to Local Ethical Research Panel approval and was conducted in accordance with the UK Animals (Scientific Procedures) Act, 1986; every effort was made to minimise pain and discomfort experienced by animals.

Electrophysiology

Preparation of acute cerebellar slices. Breeding pairs of *+/du^{2J}* mice (C57Bl/6 background) were originally supplied by Professor Annette Dolphin (University College London, UK) from which progeny were bred in-house at the University of Reading and whose genetic classification was determined by the Sequencing & Genotyping Facility, University College London from ear-notch tissue samples. Acute cerebellar brain slices were prepared from 3- to 5-week-old male mice as previously described (Ma *et al.* 2008). Briefly, animals were killed by a Schedule 1 method followed by immediate decapitation. The brain was then rapidly removed and submerged in cold, sucrose-based aCSF solution (sucrose 218 mM, KCl 3 mM, NaHCO₃ 26 mM, NaH₂PO₄ 2.5 mM, MgSO₄ 2 mM, CaCl₂ 2 mM and D-glucose 10 mM) and 300 μ m thick parasagittal cerebellar slices were prepared using a Vibroslice 725M (Campden Instruments Ltd, UK) or a Vibratome (R. & L. Slaughter, Upminster, UK). Slices were maintained under carboxygenated (95% O₂–5% CO₂), standard aCSF (NaCl 124 mM, KCl 3 mM, NaHCO₃ 26 mM, NaH₂PO₄ 2.5 mM, MgSO₄ 2 mM, CaCl₂ 2 mM and D-glucose 10 mM) at 37°C for <1 h before being returned to room temperature (22–24°C). Recordings were made at 22–24°C, 2–8 h following slice preparation.

Multi-electrode array (MEA) recording. Spontaneous unit and multi-unit spikes were recorded from acute

cerebellar slices with respect to a reference ground electrode using substrate integrated MEAs (Multi Channel Systems, Reutlingen, Germany) that consisted of 59 recording electrodes (30 μm diameter; 200 μm spacing) arranged in an 8×8 matrix minus corner electrodes, as previously described (Ma *et al.* 2008). Briefly, slices were adhered to the MEA surface and imaged via a Mikro-Okular camera (Bresser, Germany); once placed, the slice was submerged in carboxygenated standard aCSF, maintained at 24°C and perfused at a rate of $\sim 2 \text{ ml min}^{-1}$ and allowed to equilibrate for at least 15 min prior to recordings. Signals were amplified (1100 \times gain) and high-pass filtered (10 Hz) by a 60-channel amplifier (MEA60 System, MultiChannel Systems) and each channel simultaneously sampled at 10 kHz. Continuous recordings from each channel were made using MC_Rack software (MultiChannel Systems) where control spontaneous neuronal activity was first recorded for ≥ 10 min. In all experiments, each drug was bath-applied for ≥ 25 min to achieve steady-state effects before 300 s duration continuous recordings were taken. Spike events within continuous recordings were identified using MC_Rack by threshold detection at $4.5 \times$ the standard deviation of the mean of a signal-free recording. All analyses included all detected spike events that occurred during the 300 s recording period. Individual spike timings were defined by the time at which the peak minimum for each spike occurred. Spike cut outs were taken for the period 1 ms prior to and 2 ms following each spike's peak minimum (Fig. 1Aa). Spike timings were exported to Neuroexplorer4 (Nex Technologies, USA) for analysis of spike firing rates. Mean spike amplitudes were determined from spike cutout data analysed using in-house code for MATLAB 7.1 (MathWorks, Natick, MA, USA). Regularity of firing was estimated using the coefficient of variation (CV) of interspike interval (ISI), where $\text{CV} = \text{standard deviation}/\text{mean}$ and increases in CV reflect increases in firing irregularity. MEAs have previously been shown to be well suited to recording single unit activity from acute, cerebellar slices (Egert *et al.* 2002); the validity of such recordings was routinely confirmed via per electrode autocorrelograms that reliably revealed troughs at $t = 0$ s in PCs, indicative of single units. Stated replicates undertaken in MEA experiments represent the mean of electrodes for a given cellular population per slice as our unit measurement. Thus, for each slice, measured parameters (firing frequency, spike amplitude, CV) from a particular cell type were calculated for each electrode before averaging to provide a single value per cell type for a given slice. To avoid sampling bias, ≥ 6 separate slices were used for each treatment group. These data were normally distributed ($P < 0.05$, D'Agostino and Pearson omnibus normality test). Given the slice-to-slice variability in activity under control conditions, drug effects were normalised by expression of change *versus*

the starting control for each slice. Comparisons between raw measures obtained from wild-type $+/+$, $+/du^{2J}$ and du^{2J}/du^{2J} mice were performed using one-way analysis of variance followed by Tukey's HSD test or Kruskal–Wallace with Dunn's *post hoc* test as appropriate. Comparisons between multiple treatment groups were performed using Friedman's test followed by Dunn's *post hoc* test. Throughout, all data are expressed as mean \pm SEM unless stated and differences considered significant if $P < 0.05$.

Patch-clamp recording. Individual cerebellar brain slices were placed in a recording chamber maintained at room temperature and superfused with carboxygenated standard aCSF. PCs were identified morphologically using an IR-DIC upright Olympus BX50WI microscope (Olympus, Tokyo, Japan) with a $60 \times$ numerical aperture 0.9, water immersion lens. Whole-cell patch-clamp recordings from PCs were made in voltage-clamp mode with an EPC-9 patch-clamp amplifier (HEKA Elektronik, Lambrecht, Germany) using Pulse software (HEKA) on a Macintosh G4 computer (Apple Computer, Cupertino, CA, USA). Electrodes were fabricated from borosilicate glass (GC150-F10, Harvard Apparatus, Kent, UK) and had resistances $\sim 5\text{--}7 \text{ M}\Omega$ when filled with an intracellular solution (CsCl 140 mM, MgCl₂ 1 mM, CaCl₂ 1 mM, EGTA 10 mM, MgATP 4 mM, NaGTP 0.4 mM and Hepes 10 mM, pH 7.3). Series resistance was measured at 15–20 M Ω with 70–90% compensation. sIPSCs were isolated at IN-PC synapses in the presence of the non-selective ionotropic glutamate receptor antagonist, NBQX (5 μM), at a holding potential of -70 mV (Stephens *et al.* 2001). Data were sampled at 5 kHz and filtered at one-third of the sampling frequency. Drugs were diluted in aCSF and superfused ≥ 25 min and at least 150 s recording obtained during the steady-state period was used as raw data for event detection.

Data were initially exported using Pulsefit (HEKA) to AxoGraph 4.0 software for event detection using a sliding template function. Data were normally distributed ($P < 0.05$, D'Agostino and Pearson omnibus normality test). Comparisons between measures obtained from $+/+$, $+/du^{2J}$ and du^{2J}/du^{2J} mice were performed using a one-way ANOVA test followed by Tukey's HSD test. Comparison of multiple treatment groups was performed using repeated measurement one-way ANOVA, followed by Tukey's HSD test.

Radioligand binding assays

Membrane preparation. Cerebellar tissue was dissected from $+/+$, $+/du^{2J}$ or du^{2J}/du^{2J} mice (3- to 5-week-old, male) and stored separately at -80°C until use, as previously described (Jones *et al.* 2010). Tissue was suspended in a membrane buffer containing Tris-HCl 50 mM, MgCl₂ 5 mM, EDTA 2 mM and 0.5 mg ml⁻¹ fatty acid-free BSA

and complete protease inhibitor (pH 7.4, Sigma, UK) and subsequently homogenised using an Ultra-Turrax blender (IKA, UK). Homogenates were centrifuged at 1200g for 10 min and supernatants decanted. Resulting pellets were homogenised and centrifugation repeated. Pooled supernatants were then centrifuged at 39,000g for 30 min in a high-speed centrifuge (Sorvall, UK) and supernatants discarded. Remaining pellets were resuspended in membrane buffer and protein content determined by Lowry assay (Lowry *et al.* 1951).

Saturation binding assay. An initial saturation binding assay was carried out using increasing concentrations of the tritiated CB₁R antagonist, [³H]SR141716A; the CB₁R antagonist, AM251, was used as the non-specific competitor (as previously described in Jones *et al.* 2010). All concentrations tested were performed in triplicate in assay buffer (20 mM Hepes, 1 mM EDTA, 1 mM EGTA, 0.5% w/v fatty acid-free BSA, pH 7.4). All drug stocks and membrane preparations were diluted in assay buffer and stored on ice immediately prior to use. Assay tubes contained a final volume of 1 ml with [³H]SR141716A at final concentrations of (in nM): 0.1, 0.2, 0.5, 1, 2, 5, 10, 20 and a final concentration of 10 μM AM251 to determine non-specific binding. Assays were initiated by addition of 30 μg membrane protein and were incubated for 1.5 h at 25°C for ligands to reach equilibrium and terminated by rapid filtration through Whatman GF/C filters using a Brandell cell harvester. This was followed by four washes with 3 ml ice-cold PBS (0.14 M NaCl, 3 mM KCl, 1.5 mM KH₂PO₄, 5 mM Na₂HPO₄; pH 7.4) to remove unbound radioactivity. Filters were soaked in 2 ml scintillation fluid overnight. Radioactivity was quantified by liquid scintillation spectrometry using a Wallac 1414 scintillation counter where radioactivity bound to cerebellar membrane was quantified in dpm before conversion to pmol mg⁻¹.

Analyses of saturation binding assay data were conducted by non-linear regression and fitted to a one-binding site model (Jones *et al.* 2010) to determine the equilibrium dissociation constant K_d (nM) and maximal number of binding sites B_{max} (pmol mg⁻¹) using GraphPad Prism software (version 4.03; GraphPad Software Inc., San Diego, CA, USA). One-way ANOVA was used to compare results obtained from +/+, +/*du*^{2j} and *du*^{2j}/*du*^{2j} mouse tissues, followed by Tukey's HSD test when appropriate.

Pharmacology

NBQX, WIN55,212-2 (WIN55; each made up as 1000× stocks) and AM251 (made up as a 5000× stock) were dissolved in DMSO and stored at -20°C. Drug stock solutions were diluted to final desired bath concentration

using carboxygenated standard aCSF immediately before application.

Results

We have investigated the effects of *du*^{2j} mutation on cerebellar function by comparing +/+ wild-type litter-matched controls with heterozygous +/*du*^{2j}, which have >50% reduction in α2δ-2 protein (Donato *et al.* 2006), and *du*^{2j}/*du*^{2j} mice, which exhibit complete α2δ-2 ablation, reduced whole-cell PC Ca²⁺ current, an ataxic phenotype and fail to survive to adulthood (Donato *et al.* 2006).

du^{2j} mutation affects spontaneous neuronal spike activity in the cerebellum

The cerebellum consists of the PCL, whose principal PC cells represent the sole output of the cerebellar cortex, the GCL and the molecular layer that, together, provide a well-defined architecture for acute brain slice investigations of spatio-temporal network activity using multi-electrode methods (Egert *et al.* 2002; Ma *et al.* 2008). Within the PCL, *du*^{2j} mutation significantly increased spike firing irregularity in *du*^{2j}/*du*^{2j} compared with +/+ and +/*du*^{2j} mice (both $P < 0.001$; Fig. 1*Ad*); PCL spike firing frequency (Fig. 1*Ab*) and spike amplitude (Fig. 1*Ac*) were unaffected. Within the GCL, *du*^{2j}/*du*^{2j} exhibited significantly more irregular firing compared to either +/+ or +/*du*^{2j} mice (both $P < 0.01$; Fig. 1*Bd*), with no genotype-specific difference in firing frequency (Fig. 1*Bb*) or spike amplitude (Fig. 1*Bc*). Overall, these initial results reveal changes in spontaneous network firing properties resulting from *du*^{2j} mutation that most clearly manifest as globally increased PCL firing irregularity in homozygous *du*^{2j}/*du*^{2j} mice, and suggest that the major effect of *du*^{2j} mutation was to reduce cerebellar PC firing precision, potentially with secondary effects on GCL firing, an effect requiring two *du*^{2j} alleles to manifest fully.

du^{2j} mutation attenuates CB₁R modulation of spontaneous neuronal spike activity in the cerebellum

CB₁Rs are highly expressed in the cerebellum (Tsou *et al.* 1997), where they strongly regulate PC network activity and, consequentially, modulate the final output of the cerebellar cortex (Ma *et al.* 2008). Modulation of CB₁R function can cause severe motor incoordination (including ataxia), as associated with cerebellar dysfunction (DeSanty & Dar, 2001; Patel & Hillard, 2001). CB₁R modulation has been suggested as a precipitating factor for cerebellar ataxias (Smith & Dar, 2006). Therefore, we next examined CB₁R ligand

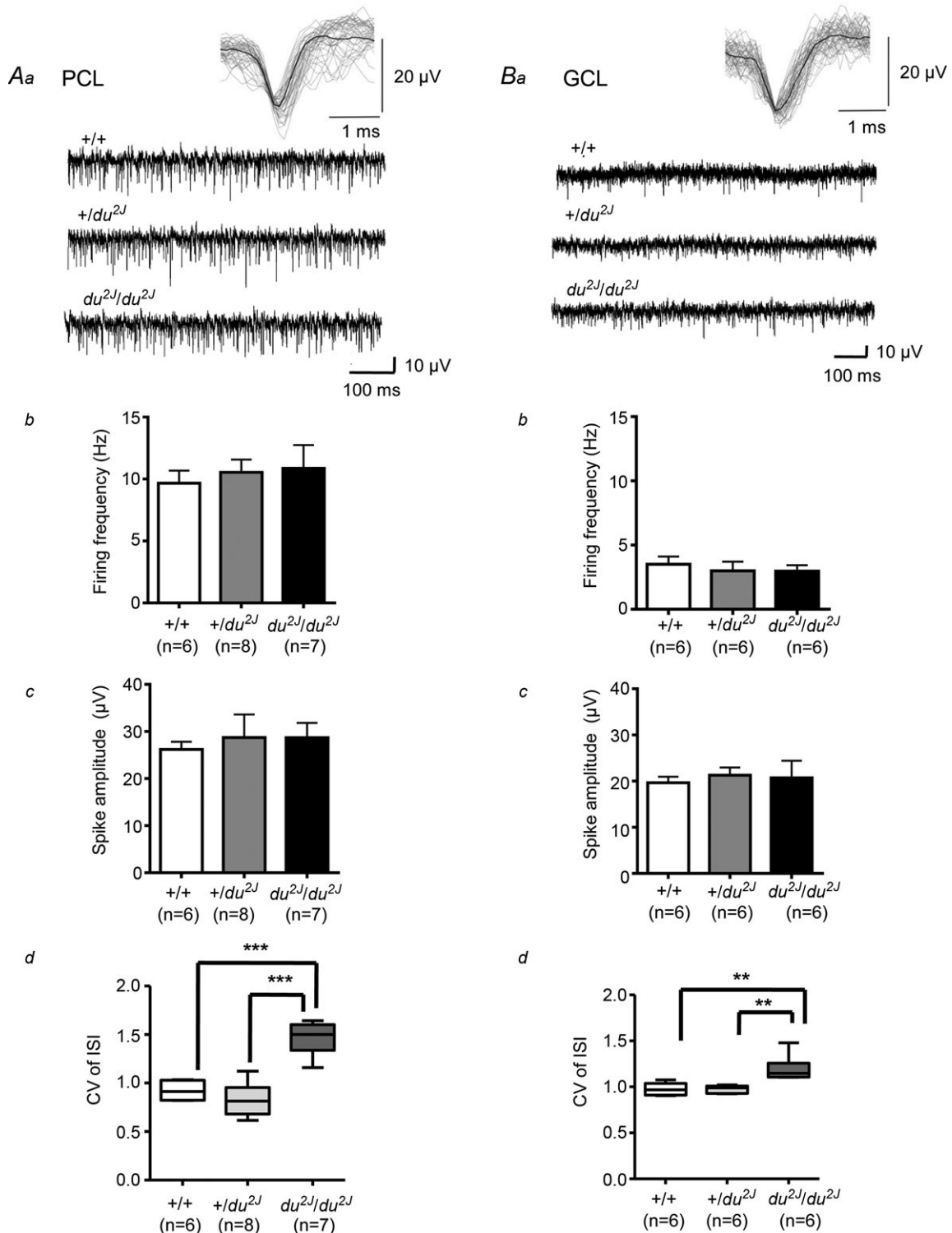


Figure 1. Region-specific comparison of basal spontaneous spike firing properties in *du^{2J}* mutants
Aa, sample traces of continuous MEA recording from a single electrode in PCL in +/+ and *du^{2J}* mutants where inset shows overlay plot of 50 spikes (grey) and mean spike shape (black) from +/+. *Ab–d*, summary bar graphs of spike firing frequency (*b*), spike amplitude (*c*) and coefficient of variation of interspike interval (CV of ISI; *d*). *du^{2J}/du^{2J}* firing was more irregular compared with +/+ and +/*du^{2J}*. *Ba*, sample traces of continuous MEA recording from a single electrode in GCL in +/+ and *du^{2J}* mutants where inset shows overlay plot of 50 spikes (grey) and mean spike shape (black) from +/+. *Bb–d*, summary bar graphs of spike frequency (*b*), spike amplitude (*c*) and CV of ISI (*d*). *du^{2J}/du^{2J}* firing was more irregular compared with +/+ and +/*du^{2J}*. ***P* < 0.01; ****P* < 0.001; Kruskal–Wallis test followed by Dunn’s test.

effects upon spontaneous neuronal activity in cerebellar slices from $+/+$, $+/du^{2f}$ and du^{2f}/du^{2f} mice. We first recapitulated our previous study (performed in TO strain mice, Ma *et al.* 2008) to confirm that the CB₁R agonist, WIN55 (5 μ M), significantly increased PCL spike firing frequency ($P < 0.05$ vs. control), an action fully reversed by subsequent application of CB₁R antagonist, AM251 (2 μ M), in the continued presence of WIN55 in $+/+$ ($P < 0.01$ vs. WIN55 only; Fig. 2Aa and b). In these experiments, neither WIN55 nor AM251 affected PCL spike amplitude (Fig. 2Ac) or spike firing regularity (Fig. 2Ad). We next investigated whether du^{2f} mutation consequentially affected CB₁R-mediated modulation of PC firing. Importantly, WIN55 (5 μ M) and AM251 (2 μ M) failed to affect PCL spike firing frequency, spike amplitude or regularity of firing in $+/du^{2f}$ (Fig. 2Ba–d) or du^{2f}/du^{2f} mice (Fig. 2Ca–d).

We next examined CB₁R ligand effects on GCL spontaneous spike firing in the du^{2f} genotypes. In $+/+$, WIN55 (5 μ M) and subsequent AM251 (2 μ M) application in the continued presence of WIN55 had no effect on GCL firing frequency (Fig. 3Aa and b), spike amplitude (Fig. 3Ac) or firing regularity (Fig. 3Ad). Similarly, WIN55 and AM251 did not affect GCL spike firing in $+/du^{2f}$ (Fig. 3Ba–d) or du^{2f}/du^{2f} mice (Fig. 3Ca–d). These results most likely reflect the reported lack of CB₁R expression in GC neurones (Tsou *et al.* 1997; Egertova & Elphick, 2000). Overall, these findings show that that CB₁R ligands predictably modulate cerebellar PCL network level activity in $+/+$, but not $+/du^{2f}$ or du^{2f}/du^{2f} , mice and are without effect on GCL firing, independent of genotype.

du^{2f} mutation affects CB₁R-mediated presynaptic inhibition at inhibitory IN-PC synapses

We have previously shown that PC firing can be affected by CB₁R-mediated modulation of presynaptic GABA release at IN-PC synapses (Ma *et al.* 2008). Given our data showing that CB₁R-modulation of spontaneous neuronal firing is absent in du^{2f} mutants, we next investigated whether du^{2f} mutation affected CB₁R modulation of inhibitory transmission at IN-PC synapses. Presynaptic Ca²⁺ channels (predominantly Ca_v2.1) underlie GABA release at IN-PC synapses (Forti *et al.* 2000; Stephens *et al.* 2001; Lonchamp *et al.* 2009) and the du^{2f} mutation has been shown to impair PC Ca²⁺ channel function (Donato *et al.* 2006). Therefore, we recorded sIPSCs to allow us to determine the effects of du^{2f} mutation on action potential-induced, Ca²⁺-mediated vesicular neurotransmitter release (Stephens *et al.* 2001) and, also, to investigate potential associations between effects at IN-PC synapses and the action potential-dependent spontaneous PC spike firing measurements described above. No significant differences in sIPSC frequency (Fig. 4A and Ba)

or regularity (Fig. 4A and Bc) between $+/+$, $+/du^{2f}$ and du^{2f}/du^{2f} were observed, although $+/du^{2f}$ and du^{2f}/du^{2f} each exhibited significantly increased sIPSC amplitudes when compared with $+/+$ mice ($+/du^{2f}$: $P < 0.05$; du^{2f}/du^{2f} : $P < 0.01$; Fig. 4Bb).

We next confirmed the predicted CB₁R modulation of sIPSC frequency at $+/+$ IN-PC synapses (Takahashi & Linden, 2000; Szabo *et al.* 2004). Thus, WIN55 (5 μ M) significantly decreased sIPSC frequency ($P < 0.05$), an effect that was reversed and increased beyond control levels by subsequent AM251 (2 μ M) application in $+/+$ mice ($P < 0.01$; Fig. 5Aa and b). The latter result is consistent with the presence of endocannabinergic tone or constitutive CB₁R activity in this system (Ma *et al.* 2008; Wang *et al.* 2011). Consistent with the lack of CB₁R-mediated effects on neuronal spiking activity described above, WIN55 and AM251 failed to significantly modulate sIPSC frequency in $+/du^{2f}$ (Fig. 5Ba and b) or du^{2f}/du^{2f} (Fig. 5Ca and b), although both WIN55 and AM251 showed a marginal trend ($P = 0.07$; repeated measurement one-way ANOVA) to modulate sIPSC frequency in $+/du^{2f}$ (Fig. 5Bb), an effect not seen in du^{2f}/du^{2f} ($P = 0.19$; Fig. 5Cb). In addition, WIN55 significantly increased sIPSC amplitude in $+/+$ ($P < 0.05$; Fig. 5Ac) and $+/du^{2f}$ ($P < 0.05$; Fig. 5Bc), but not du^{2f}/du^{2f} ($P = 0.11$; Fig. 5Cb). Subsequent AM251 application was without effect on WIN55-induced increases in sIPSC amplitude in $+/+$ (Fig. 5Ac), $+/du^{2f}$ (Fig. 5Bc) or du^{2f}/du^{2f} mice (Fig. 5Cc). The inability of AM251 to block WIN55-induced increases in sIPSC amplitude suggests a CB₁R-independent action here.

Taken together, these results demonstrate an attenuation of CB₁R modulation at IN-PC synapses in du^{2f} mutants. Such effects could contribute to the observed deficits in network level neuronal function.

Investigation of CB₁ receptor expression in du^{2f} mice using [³H]SR141716A saturation binding assay

The data above demonstrate that du^{2f} mutants exhibit deficits in CB₁R-mediated signalling in the cerebellum. Such deficits could occur as a consequence of reported defects in $\alpha 2\delta$ -2 Ca²⁺ channel subunit expression (Donato *et al.* 2006); however, an alternative hypothesis is reduced CB₁R expression in the cerebella of du^{2f} mutants. To further investigate the latter hypothesis, CB₁R expression was investigated using radioligand saturation binding assays. In $+/+$, $+/du^{2f}$ and du^{2f}/du^{2f} mice, specific binding of the high-affinity CB₁R antagonist [³H]SR141716A to cerebellar membranes was concentration dependent and saturable (Fig. 6). There was no significant difference in K_d between $+/+$, $+/du^{2f}$ and du^{2f}/du^{2f} ($P = 0.47$; Table 1) and the Hill coefficient (n_H , the gradient of the Hill plot)

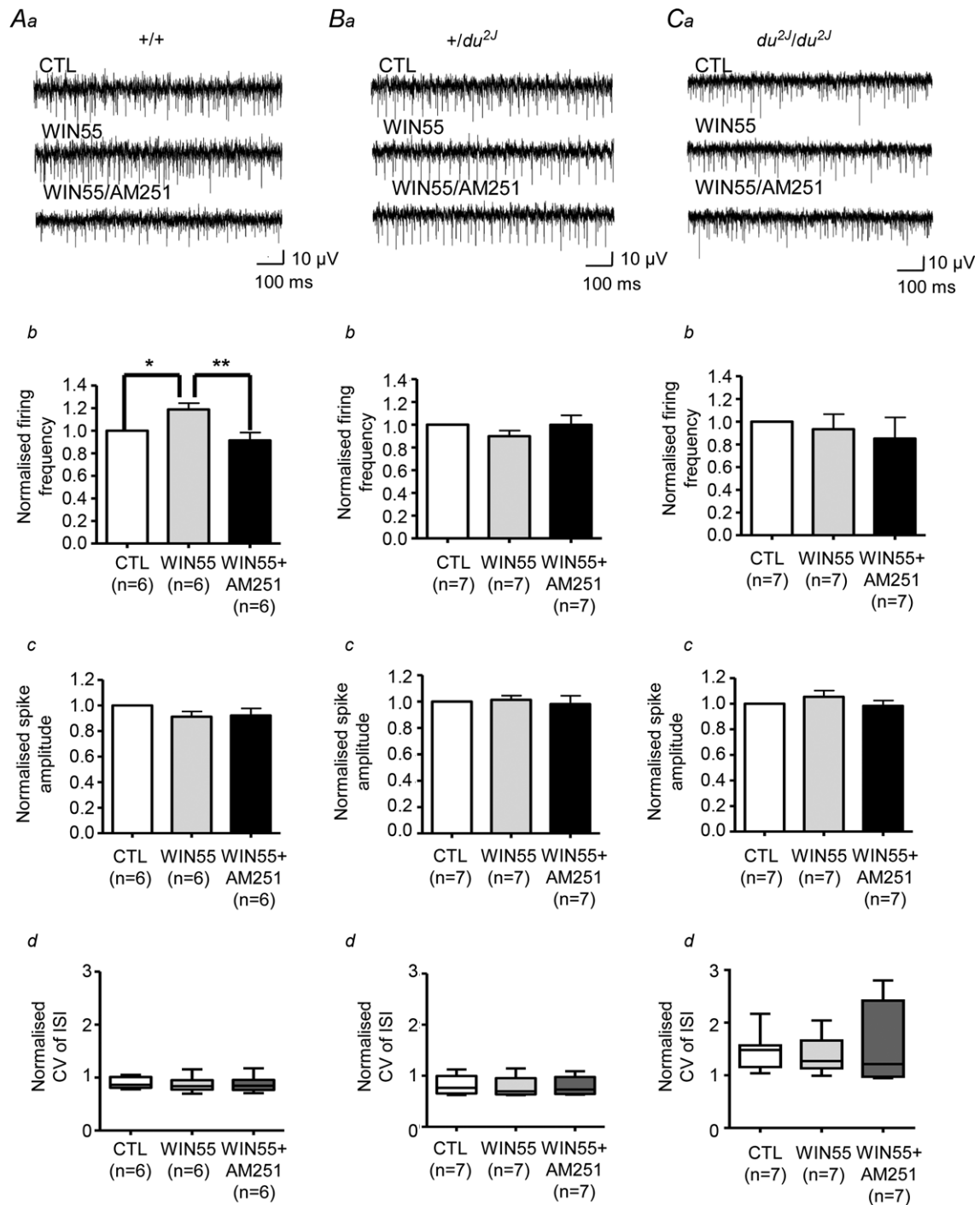


Figure 2. Differential effects of CB₁R ligands on spontaneous PCL spike activity in *du^{2J}* mutants
 Aa–Ca, sample traces of continuous MEA recording from a single electrode in PCL showing effect of WIN55 (5 μM) and subsequent application of AM251 (2 μM; in the continued presence of 5 μM WIN55) on spontaneous spike firing in +/+ (Aa), +/*du^{2J}* (Ba) and *du^{2J}/du^{2J}* (Ca). Ab–Cd, summary bar graphs showing that WIN55 significantly increased normalised spike firing frequency and subsequent application of AM251 caused a significant decrease in normalised spike firing frequency in +/+ (Ab). By contrast, WIN55 and subsequent application of AM251 had no significant effect on normalised spike firing frequency in +/*du^{2J}* (Bb) or *du^{2J}/du^{2J}* (Cb). CB₁R ligands had no effect on spike amplitude in +/+ (Ac), +/*du^{2J}* (Bc) or *du^{2J}/du^{2J}* (Cc) or on normalised CV of ISI in +/+ (Ad), +/*du^{2J}* (Bd) or *du^{2J}/du^{2J}* (Cd). **P* < 0.05; ***P* < 0.01; Friedman test followed by Dunn’s test.

approximated unity for all genotypes (Table 1), indicating that [^3H]SR141716A bound at a single site to cerebellar CB₁Rs. Most importantly, cerebellar membranes from +/+, +/*du*^{2J} and *du*^{2J}/*du*^{2J} mice exhibited no significant differences in B_{max} ($P = 0.3$; Table 1), indicating that there

was no difference in CB₁R expression between genotypes investigated. These data demonstrate that the reported deficit in CB₁R signalling in *du*^{2J} mutants was likely not to be due to reduced CB₁R expression and, rather, may reflect defects in $\alpha 2\delta$ -2 expression, as discussed below.

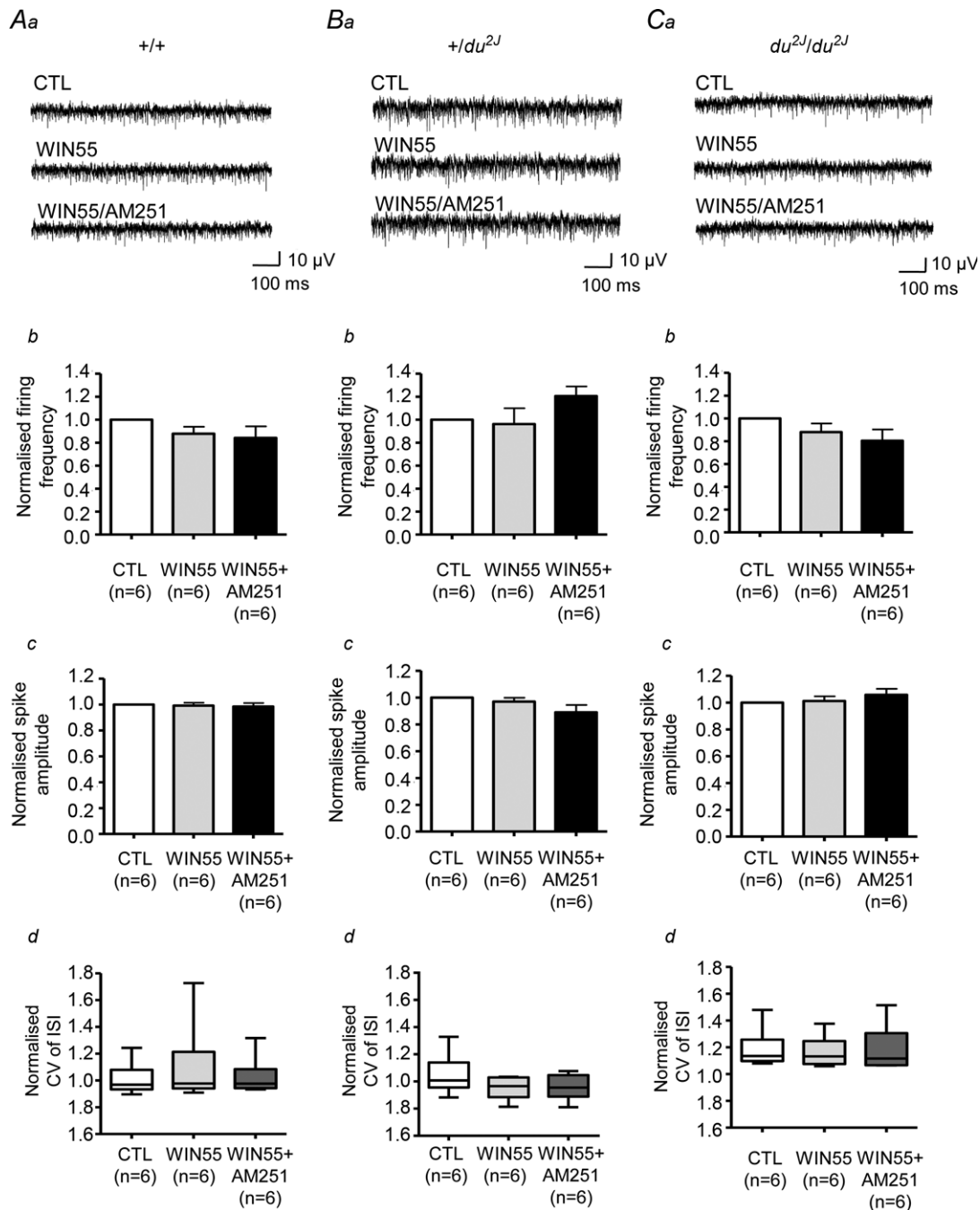


Figure 3. Lack of effect of CB₁R ligands on spontaneous GCL spike activity in *du*^{2J} mutants

Aa–Ca, sample traces of continuous MEA recording from a single electrode in GCL showing lack of effect of WIN55 (5 μM) and subsequent application of AM251 (2 μM); in the continued presence of 5 μM WIN55) on spontaneous spike firing in +/+ (Aa), +/*du*^{2J} (Ba) and *du*^{2J}/*du*^{2J} (Ca). Ab–Cd, summary bar graphs showing that CB₁R ligands had no effect on normalised spike firing frequency in +/+ (Ab), +/*du*^{2J} (Bb) or *du*^{2J}/*du*^{2J} (Cb), or on spike amplitude in +/+ (Ac), +/*du*^{2J} (Bc) or *du*^{2J}/*du*^{2J} (Cc), or on normalised CV of ISI in +/+ (Ad), +/*du*^{2J} (Bd) or *du*^{2J}/*du*^{2J} (Cd); as assessed by Friedman test followed by Dunn's test.

Discussion

$\alpha 2\delta$ -2 mouse mutants exhibit ataxia. Here, we use the *du^{2J}* mutation, a reportedly clean $\alpha 2\delta$ -2 knockout (Donato *et al.* 2006) permitting clear interpretation of phenotypic differences. In addition to studying homozygous *du^{2J}/du^{2J}* mice, we also examine heterozygous *+/du^{2J}* mice to investigate potential progressive disturbances. We demonstrate that *du^{2J}* mutants exhibit deficits in cerebellar CB₁R-mediated signalling.

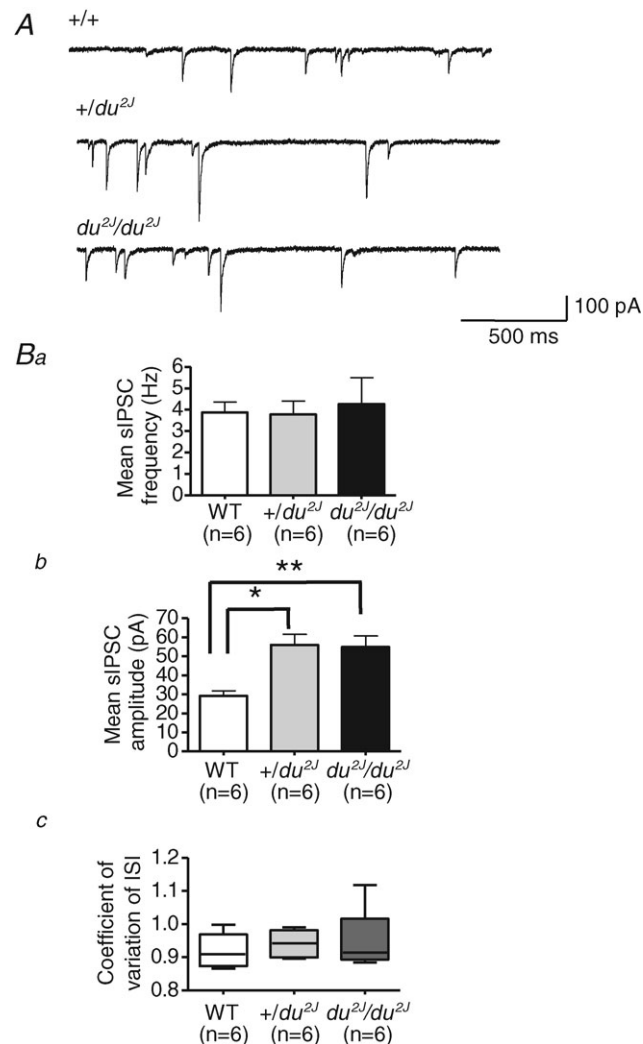


Figure 4. Comparison of basal spontaneous inhibitory transmission at IN-PC synapses in *du^{2J}* mutants

A, raw sIPSC traces from representative PCs from *+/+*, *+/du^{2J}* and *du^{2J}/du^{2J}*. Ba–c, summary bar graphs showing that there was no significant differences in mean sIPSC frequency (a) and CV of ISI (c), but that mean sIPSC amplitude (b) was significant increased in *+/du^{2J}* and *du^{2J}/du^{2J}* compared to *+/+*. * $P < 0.05$; ** $P < 0.01$; one-way analysis of variance followed by Tukey's HSD test.

Effects of *du^{2J}* mutation on neuronal spike activity in the cerebellum

PCL and GCL spike firing showed negative polarity (Egert *et al.* 2002; Ma *et al.* 2008). PCL spikes on a given electrode arose from single cells as supported by characteristic trough autocorrelograms and single distribution ISI histograms (data not shown). Conversely, GCL spikes produced variable distribution ISI histograms and autocorrelograms that suggested multi-cell signals (data not shown), which can be accounted for by larger cell somata diameters in PCL than GCL (Egert *et al.* 2002). GCL spike recordings using MEAs show some differences in the literature, ranging from reports of regular activity consistent with the present findings (Egert *et al.* 2002), to recordings that are 'usually silent' and where the sparse spontaneous activity seen was attributed to Golgi cell activity (Mapelli & D'Angelo, 2007). However, care should be taken when making comparisons between reports where experimental conditions vary (e.g. recordings made at 22–24°C here and in Egert *et al.* (2002) vs. 32°C in Mapelli & D'Angelo (2007), which can have a profound effect upon basic firing properties. The above caveats for GCL notwithstanding, the major effect of the *du^{2J}* mutation was to increase PCL irregularity without affecting firing frequency or spike amplitude. *du^{2J}/du^{2J}*, but not *+/du^{2J}*, exhibited increased PC firing irregularity suggesting progressive dysfunction; this effect could be coupled to differential reduction of $\alpha 2\delta$ -2 protein expression between *+/du^{2J}* and *du^{2J}/du^{2J}* mice (50% vs. 100% respectively, Donato *et al.* 2006). We confirm that expression of two *du^{2J}* alleles is required for manifestation of increased PC irregularity and an ataxic phenotype. Although *du^{2J}* cerebella are smaller than *+/+*, *du^{2J}* mutants show no differences in dendritic morphology (Donato *et al.* 2006), arguing against PC degeneration underlying differences in firing regularity. Both PC firing precision and activity patterns play important roles in cerebellar motor control (Womack & Khodakhah, 2002; De Zeeuw *et al.* 2011), potentially by time-locking PC spiking activity (Person & Raman, 2011). Such precision is affected by behavioural state and tactile stimulation (Shin *et al.* 2007). Importantly, many Ca²⁺ channel mutants, including *du* and *du^{2J}*, increase PC firing irregularity (Hoebeek *et al.* 2005; Donato *et al.* 2006; Walter *et al.* 2006; Ovsepian & Friel, 2012; Alviña & Khodakhah, 2010), which is predicted to adversely affect cerebellar function; for example, PC firing irregularity in *tottering* mutants functionally reduces compensatory eye movement amplitude (Hoebeek *et al.* 2005). Donato *et al.* (2006) reported reduced spontaneous PC firing frequency in *+/du^{2J}* that was further reduced in *du^{2J}/du^{2J}*, although this was not observed here or in studies using *du* mutants (Walter *et al.* 2006). These differences may be developmental, since younger animals were used by

Donato *et al.* (2006) in comparison to those used in our study and by Walter *et al.* (2006); however, it is clear that the major, consistent effect of the *du^{2J}* mutation is to increase firing irregularity.

It has been proposed that GCL firing, driven by mossy fibre inputs, manifests as precisely timed spike bursts limited by Golgi cell-mediated feedforward inhibition to form discrete time windows (~5 ms) for control of

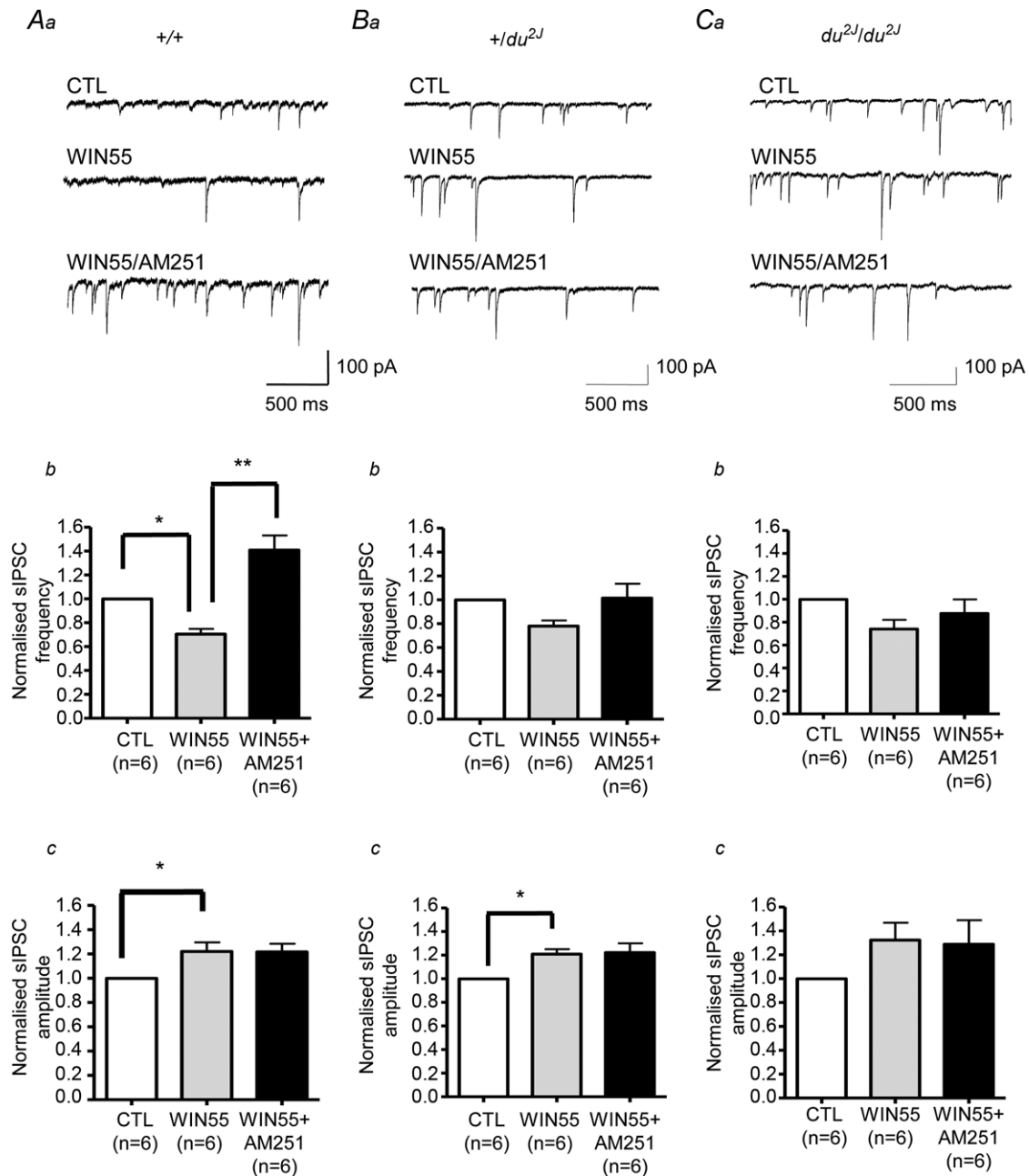


Figure 5. Differential effects of CB₁R ligands on inhibitory transmission at IN-PC synapses in *du^{2J}* mutants

Aa–Ca, raw sIPSC traces from representative PCs from +/+ (Aa), +/*du^{2J}* (Ba) and *du^{2J}/du^{2J}* (Ca) showing effect of WIN55 (5 μM) and subsequent application of AM251 (2 μM; in the continued presence of 5 μM WIN55). Ab–Cb, summary bar graphs showing that WIN55 significantly reduced and AM251 significantly increased normalised sIPSC frequency in +/+ (Ab), but was without effect in +/*du^{2J}* (Bb) or *du^{2J}/du^{2J}* (Cb). Ac–Cc, WIN55 significantly increased normalised sIPSC amplitude in +/+ (Ac) and +/*du^{2J}* (Bc), but was without effect in *du^{2J}/du^{2J}* (Cc). Subsequent application of AM251 was without effect in each case. **P* < 0.05; ***P* < 0.01; repeated measurement one-way ANOVA followed by Tukey's HSD test.

distinct motor domains; thus, GC spike firing dysfunction could contribute to ataxic symptoms (e.g. hypermetria; D'Angelo & De Zeeuw, 2009). Here, GCL firing irregularity was increased in *du^{2J}/du^{2J}* mice, although to a far lesser extent than in PCL. During development, GC survival depends upon connectivity with PCs (Lossi *et al.* 2002) and PC disturbances adversely affect GC (Goldowitz & Hamre, 1998), with PC-dependent GCL degeneration proposed as a mechanism (Ivanov *et al.* 2004); this phenomenon is also reported for ataxic *lurcher* mice

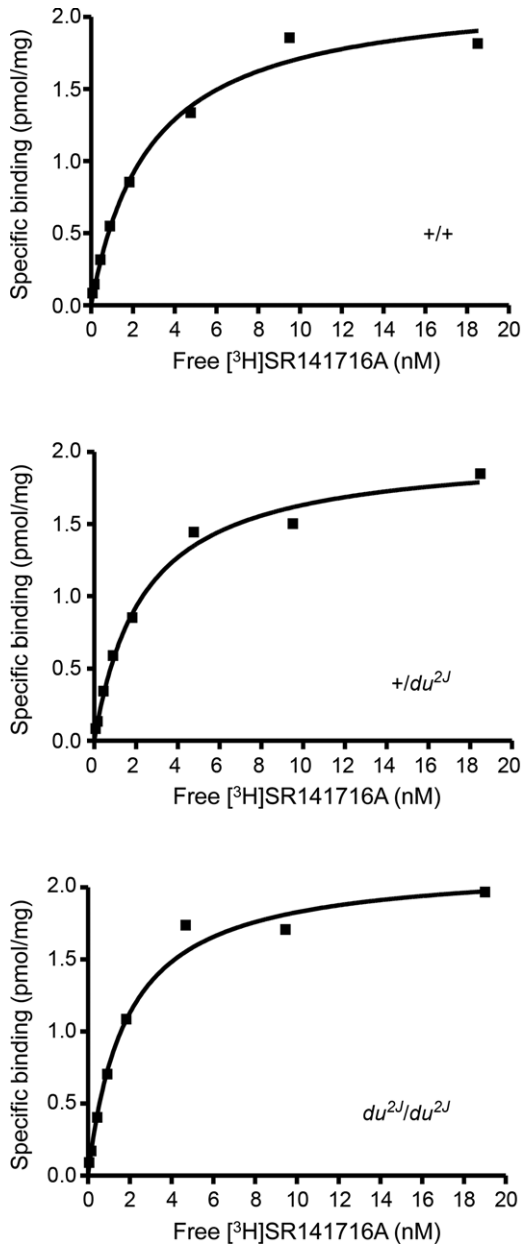


Figure 6. Saturation binding of [³H]SR141716A to cerebellar membranes in *du^{2J}* mutants

Representative saturation binding curve for [³H]SR141716A in cerebellar membranes from +/+ (A), +/*du^{2J}* (B) and *du^{2J}/du^{2J}* (C).

Table 1. [³H]SR141716A saturation binding data for cerebellar membrane in *du^{2J}* mutants

Genotype	K_d (nM)	B_{max} (pmol mg ⁻¹)	n_H
+/+ ($n = 3$)	3.1 ± 0.2	2.15 ± 0.08	0.99 ± 0.01
+/ <i>du^{2J}</i> ($n = 3$)	2.9 ± 0.3	2.34 ± 0.12	0.99 ± 0.02
<i>du^{2J}/du^{2J}</i> ($n = 4$)	2.4 ± 0.3	2.03 ± 0.21	1.01 ± 0.01

K_d and B_{max} were obtained from the saturation binding curves plotted between specific binding vs. free [³H]SR141716A radioligand concentration. No significant differences in K_d ($P = 0.47$) or B_{max} ($P = 0.3$) were seen; one-way analysis of variance. Hill slope (n_H) was obtained from the Hill plot of the data transformed from saturation binding plot

(Wetts & Herrup, 1982). Interestingly, $\alpha 2\delta$ -2 subunits are barely expressed in GCL, and GC Ca²⁺ currents were normal in *du* mutants (Barclay *et al.* 2001; Donato *et al.* 2006), consistent with GC changes reflecting secondary consequences of PC dysfunction. Overall, although connectivity deficiencies between cerebellar layers in *du^{2J}* mutants remain unproven, our results provide evidence for a role of $\alpha 2\delta$ -2 in correct PC–GC signalling and suggest that the impact of $\alpha 2\delta$ -2 loss on the GCL should not be ignored.

Effects of *du^{2J}* mutation on inhibitory synaptic transmission in the cerebellum

Whilst effects of *du^{2J}* mutation on synaptic transmission are unknown, ataxic mouse models exhibit differences in excitatory transmission in some studies (Matsushita *et al.* 2002; Liu & Friel, 2008), but not others (Zhou *et al.* 2003); *leaner* mutants exhibit enhanced inhibitory transmission, proposed to underlie reduced PC firing and increased irregularity (Liu & Friel, 2008). In addition to intrinsic properties, tonic inhibitory inputs also regulate PC output and synchronization (Hausser & Clark, 1997; de Solages *et al.* 2008). Here, sIPSC frequencies were unaffected between genotypes, suggesting that action potential-mediated, basal GABA release is unaltered by *du^{2J}* mutation. Interestingly, sIPSC amplitude was significantly increased in +/*du^{2J}* and *du^{2J}/du^{2J}* (cf. +/+ littermates). A similar increase has been reported for *leaner* mutants and attributed to increased presynaptic GABA release (Ovsepian & Friel, 2012); such effects are unlikely here due to the reported lack of change to sIPSC frequency. An alternative hypothesis is that the mutation leads to an increase in postsynaptic GABA_A receptor responsiveness. Increased intracellular Ca²⁺ ([Ca²⁺]_i) can suppress postsynaptic GABA_A receptor function, potentially by decreasing GABA affinity for GABA_A receptors (Inoue *et al.* 1986; Martina *et al.* 1994); therefore, reduced [Ca²⁺]_i, resulting from decreased PC $\alpha 2\delta$ -2 expression in *du^{2J}* mutants (Donato *et al.* 2006), may

relieve Ca^{2+} -mediated suppression of GABA_A receptor function.

CB₁R modulation is abolished in *du*^{2J} mutants

Whilst we found no changes to basal IN-PC inhibitory transmission, it remains possible that *du*^{2J} mutation disrupts presynaptic regulatory mechanisms, including GPCR-mediated inhibition (Zhou *et al.* 2003). Here, no CB₁R-mediated modulation was seen in *+/du*^{2J} and *du*^{2J}/*du*^{2J} mice, as demonstrated by an absence of CB₁R agonist-mediated increases in PC spike firing and no reduction in inhibitory transmission at IN-PC synapses compared to *+/+* mice. These findings suggest that deficits in CB₁R presynaptic inhibition of GABA release are associated with this model of ataxia and could contribute to compromised normal regulation of total PC output and, potentially, the aberrant motor phenotype associated with deficient PC function.

Unlike changes to PC firing regularity, which were confined to homozygous *du*^{2J}/*du*^{2J} mice, heterozygous *+/du*^{2J} mice showed CB₁R signalling deficits similar to *du*^{2J}/*du*^{2J} mutants. However, WIN55 and AM251 showed a statistical trend ($P < 0.1$) to modulate sIPSC frequency in *+/du*^{2J} not seen in *du*^{2J}/*du*^{2J} mice, offering some support for a progressive deficit in modulation of presynaptic inhibition. Somewhat unexpectedly, WIN55 increased sIPSC amplitude in *+/+* and *+/du*^{2J} mice, and this increase may reflect a postsynaptic phenomenon; in this regard, the lack of AM251-induced reversal of WIN55 effects (*cf.* Wang *et al.* 2011) suggests that this WIN55 effect is CB₁R independent, consistent with the reported lack of postsynaptic CB₁R expression (Tsou *et al.* 1997; Yamasaki *et al.* 2006). For example, WIN55 has been shown to inhibit Ca_v2.1 channels in PCs at concentrations used here (Fisyunov *et al.* 2006; Lozovaya *et al.* 2009), and such actions could reduce $[\text{Ca}^{2+}]_i$ to overcome Ca^{2+} -mediated suppression of GABA_A receptor function (Inoue *et al.* 1986; Martina *et al.* 1994) in *+/+* and *+/du*^{2J} mice; the lack of effect in *du*^{2J}/*du*^{2J} mice may reflect reduced PC Ca^{2+} current levels in homozygotes (Donato *et al.* 2006). Overall, whilst expression of two *du*^{2J} alleles is required for increased PC irregularity and ataxia, our results demonstrate that expression of a single *du*^{2J} allele compromises CB₁R signalling, prior to any measurable change in PC firing regularity and any clear ataxic phenotype. Here, disrupted cannabinergic signalling may represent a useful diagnostic biomarker of early or asymptomatic cerebellar dysfunction.

Consequences of *du*^{2J} mutation for CB₁R signalling

We show, for the first time, that $\alpha 2\delta$ -2 deficits caused by *du*^{2J} mutation are associated with aberrant CB₁R

signalling and suggest links between impaired Ca^{2+} channel function and consequential impairment of GPCR-mediated presynaptic inhibition. We also show that CB₁R expression is unchanged in *du*^{2J} mutants, suggesting that deficiency occurs downstream of receptor activation. $\alpha 2\delta$ -2 is the major isoform expressed in PCs (Cole *et al.* 2005) and reduced $\alpha 2\delta$ -2 expression in *du*^{2J} affects Ca^{2+} current levels (Donato *et al.* 2006). Moreover, $\alpha 2\delta$ -2 is predominantly associated with Ca_v2.1 (Barclay *et al.* 2001), the major Ca_v α subunit mediating presynaptic GABA release at IN-PC synapses (Stephens *et al.* 2001). Importantly, PC-specific conditional Ca_v2.1 knock-out causes cerebellar ataxia (Todorov *et al.* 2012). The association of $\alpha 2\delta$ -2 and Ca_v2.1 subunits suggests that deficits in either subunit could equally cause motor deficits, as supported by similarities in ataxic phenotypes in $\alpha 2\delta$ -2 mutants, including *du*^{2J} and Ca_v2.1 knockouts. The most parsimonious explanation for our results is that altered $\alpha 2\delta$ -2 expression in axon terminals of basket and stellate interneurons in *du*^{2J} mutants leads to deficits in CB₁R-mediated signalling. Although the expression of $\alpha 2\delta$ -2 in interneurone terminals in cerebellum has not been studied specifically, $\alpha 2\delta$ -2 is highly expressed in the molecular layer and in GABAergic interneurons throughout the CNS, as well as in PCs (Barclay *et al.* 2001; Cole *et al.* 2005). Recent studies have shown that $\alpha 2\delta$ subunits affect release properties of the Ca^{2+} channel complex at presynaptic terminals by improving spatial coupling between Ca^{2+} influx and exocytosis (Hoppa *et al.* 2012; Dolphin, 2012), in addition to protecting against block of exocytosis by intracellular Ca^{2+} chelators (Hoppa *et al.* 2012). Such findings are consistent with the hypothesis that proper $\alpha 2\delta$ -2 expression is required for correct modulation of presynaptic release. Presynaptic CB₁R activation limits transmitter release via generation of G $\beta\gamma$ subunits which inhibit Ca^{2+} channels (Twitchell *et al.* 1997; Stephens, 2009). Here, reduced $\alpha 2\delta$ -2 in *du*^{2J} mutants could alter G protein– Ca^{2+} channel interaction to limit direct effects upon channel gating and so dysfunctionally affect modulation of GABA release onto PCs.

Functional impact of CB₁R deficits in cerebellar ataxia

We propose that CB₁R signalling deficits in *du*^{2J} mutants occur as a consequence of reduced $\alpha 2\delta$ -2 expression, which impairs Ca^{2+} channel function and affects normal GPCR presynaptic inhibition in ataxic phenotypes. Under normal conditions, CB₁R inhibition of GABA release at IN-PC synapses reduces inhibitory drive onto PCs to increase PC spike firing (Ma *et al.* 2008). Regulation of PC spike firing and regularity modulates activity of deep cerebellar nuclei to control motor function. CB₁R signalling also contributes to presynaptically

expressed synaptic plasticity in the cerebellar cortex. Whilst long-term depression of transmitter release is typically associated with the excitatory parallel fibre (PF)–PC pathway, endocannabinoid-mediated short term plasticity, in the form of depolarization-induced suppression of inhibition, is prominent at IN-PC synapses (Kano *et al.* 2009). Notably, CB₁R immunoreactivity is reportedly five times higher at IN than at PF terminals; in particular, at basket cell terminals at the PC axon initial segment (Kawamura *et al.* 2006). Therefore, deficits in CB₁R signalling may directly influence PC output in ataxic phenotypes, both in terms of spike firing and regularity and also synaptic function; such deficiencies may contribute to disease.

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Additional information

Competing interests

None.

Author contributions

All the authors contributed to the conception and design of the experiments, the collection, analysis and interpretation of data, and drafting the article or revising it critically for important intellectual content. All authors approved the final version of the manuscript.

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