

Measurement of steroid concentrations in brain tissue: methodological considerations

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Matthew D. Taves, Department of Psychology, University of British Columbia, 2136 West Mall, Vancouver, BC, Canada V6T 1Z4. e-mail: taves@zoology.ubc.ca It is well recognized that steroids are synthesized de novo in the brain (neurosteroids). In addition, steroids circulating in the blood enter the brain. Steroids play numerous roles in the brain, such as influencing neural development, adult neuroplasticity, behavior, neuroinflammation, and neurodegenerative diseases such as Alzheimer's disease. In order to understand the regulation and functions of steroids in the brain, it is important to directly measure steroid concentrations in brain tissue. In this brief review, we discuss methods for the detection and quantification of steroids in the brain. We concisely present the major advantages and disadvantages of different technical approaches at various experimental stages: euthanasia, tissue collection, steroid extraction, steroid separation, and steroid measurement. We discuss, among other topics, the potential effects of anesthesia and saline perfusion prior to tissue collection; microdissection via Palkovits punch; solid phase extraction; chromatographic separation of steroids; and immunoassays and mass spectrometry for steroid quantification, particularly the use of mass spectrometry for "steroid profiling." Finally, we discuss the interpretation of local steroid concentrations, such as comparing steroid levels in brain tissue with those in the circulation (plasma vs. whole blood samples; total vs. free steroid levels). We also present reference values for a variety of steroids in different brain regions of adult rats. This brief review highlights some of the major methodological considerations at multiple experimental stages and provides a broad framework for designing studies that examine local steroid levels in the brain as well as other steroidogenic tissues, such as thymus, breast, and prostate.

Keywords: corticosterone, estradiol, mouse, songbird, testosterone

INTRODUCTION

Steroids play many important physiological roles in the brain and influence brain development, behavior, cognition, neuroplasticity, and neuroinflammation (Baulieu, 1998; Sakamoto et al., 2001b; Sorrells and Sapolsky, 2009). In addition to steroids that enter the brain from the periphery, steroids are synthesized within the brain itself ("neurosteroids"; Corpéchot et al., 1981). Neurosteroids can be synthesized de novo from cholesterol or from circulating inactive precursors (Compagnone and Mellon, 2000; Schmidt et al., 2008; Do Rego et al., 2009). Because the brain is both a producer and a target of steroids, it is of great interest and utility to measure endogenous steroid levels in specific brain regions. Measurement of local steroid levels allows for comparisons among different brain regions, as well as comparisons between brain and blood. Such comparisons are useful in determining which tissues are synthesizing or metabolizing steroids (Schmidt and Soma, 2008; Hojo et al., 2009; Konkle and McCarthy, 2011). Indeed, the presence of high pregnenolone and dehydroepiandrosterone (DHEA) concentrations in the brain, and low concentrations in the plasma, of gonadectomized and adrenalectomized rats was the first evidence that suggested steroid production in the brain (Corpéchot et al., 1981). High local steroid levels, along with studies of steroidogenic enzyme mRNA, protein, and activity in the brain, have demonstrated that the brain functions as a steroid-synthetic organ (Schmidt et al., 2008; Taves et al., 2011).

The conceptual leap that permitted the brain to be considered a source (rather than merely a target) of steroid hormones meant that measuring peripheral steroid levels in the blood was no longer sufficient for understanding steroid actions in the brain. Determining the steroid level required for induction of a particular behavior, for example, involves addressing the contributions of steroids of both peripheral and central origin. To add to the complexity, peripheral steroids that enter the brain are converted into both inactive metabolites and more potent steroids (Mellon and Griffin, 2002). Since the expression and activity of steroidogenic enzymes determines the steroidal milieu of different brain regions, understanding the distribution and regulation of steroidogenic enzymes in the central nervous system is a major

step forward in estimating local steroid concentrations (Do Rego et al., 2009). Indeed, the majority of papers on this topic have measured steroidogenic enzymes in the brain, rather than the products of these enzymes (i.e., the steroids themselves). However, the aggregate result of steroid synthesis and metabolism in the brain and steroid uptake from the blood can only be determined via empirical measurements of steroid concentrations in discrete brain regions.

Measuring steroids in brain tissue presents a number of technical challenges. These challenges include: the high lipid content of brain tissue (requiring good sample preparation to avoid assay interference; Rash et al., 1980); the high spatial heterogeneity of the brain (resulting in small tissue amounts; Charlier et al., 2010); low steroid concentrations, in some cases (Gomez-Sanchez et al., 2005); ability of steroids to freely diffuse across cell membranes (due to their hydrophobic nature); unique steroids that are absent in the periphery (e.g., 17α -estradiol; Toran-Allerand et al., 2005); and rapid changes in steroid production and metabolism (Balthazart and Ball, 2006). In this brief review, we discuss methodological considerations that address or reduce these technical challenges, and we also discuss the interpretation of tissue steroid concentrations. For a more general and comprehensive guide to steroid analysis, we direct readers to the excellent text edited by Makin and Gower (2010). Although here we focus on steroids in brain tissue, the general principles discussed are also relevant for the measurement of steroids in various other organs and tissues (Taves et al., 2011).

SAMPLE COLLECTION

STUDY SPECIES

Building upon the initial description of neurosteroid synthesis in the mammalian brain (Corpéchot et al., 1981; Le Goascogne et al., 1987), subsequent work has identified neurosteroid synthesis across vertebrate species, including fish (Forlano et al., 2001; Sakamoto et al., 2001a; Diotel et al., 2011), amphibians (Mensah-Nyagan et al., 1996; Takase et al., 1999; Matsunaga et al., 2004; Bruzzone et al., 2010), and birds (Tsutsui et al., 2000; Soma et al., 2004; London et al., 2006; Saldanha et al., 2011). The high levels of steroidogenic enzyme expression and activity (and corresponding high levels of steroids) in the brains of non-mammalian species are a major advantage in the study of neurosteroids (Goodson et al., 2005; Do Rego et al., 2009; Saldanha et al., 2009; Diotel et al., 2010). In addition, because many birds are diurnal and terrestrial, they are highly amenable to field work. This allows studies of neurosteroids in a natural and ecologically relevant context and studies of how neurosteroids regulate natural behaviors (Soma and Wingfield, 2001; Soma, 2006; Soma et al., 2008; Newman and Soma, 2009; Pradhan et al., 2010b). Comparative work has shown that neurosteroid production is conserved across vertebrates, and thus appears to be a general characteristic of the vertebrate brain (Callard et al., 1978), including the human brain (Stoffel-Wagner, 2001; Yu et al., 2002).

INDIRECT MEASUREMENT OF BRAIN STEROID LEVELS

There are several strategies to indirectly assess brain steroid levels. Some of these approaches allow for repeated sampling from individual subjects, either in different contexts or in real-time

during experiments. One method of indirectly estimating brain steroid levels involves collecting blood leaving the brain (i.e., from the jugular vein) and comparing this to blood from another source (e.g., carotid artery, brachial vein, or heart). If steroid levels are higher in blood from the jugular vein, then this suggests production of these steroids in the brain (Schlinger and Arnold, 1992, 1993; Saldanha and Schlinger, 1997; Chin et al., 2008; Newman et al., 2008b; Charlier et al., 2009; Newman and Soma, 2011). To our knowledge, this approach has only been used in studies of songbirds but might prove useful in other species as well. Clearly, this approach is indirect and cannot reveal the absolute concentrations of steroids in specific brain regions. Nonetheless, this approach can be useful for determining relative changes in whole brain steroid production, and it permits repeated sampling from subjects. However, note that stress can rapidly alter steroid concentrations in the jugular blood (Newman et al., 2008b).

Cerebrospinal fluid (CSF) can also be used to indirectly estimate the neural steroidal milieu (Guazzo et al., 1996). For example, in humans, CSF steroid levels are altered in patients with neurological diseases such as Alzheimer's disease (Kim et al., 2003). However, collecting CSF samples of sufficient volume from small animals (e.g., mice) is a more difficult obstacle.

Microdialysis allows quantification of changing steroid levels in a specific brain region (Saleh et al., 2004; Droste et al., 2008; Remage-Healey et al., 2008). Artificial CSF is perfused into the brain through a microdialysis probe, and dialysate is collected across a membrane. This extremely powerful technique can be used to measure temporal changes in steroid levels at a specific location in the brain of a freely behaving subject. Another advantage is that subjects require minimal handling at the time of sample collection. A disadvantage is that steroid levels are measured in only a single region at a time. Moreover, steroid recoveries across the dialysis membrane in vitro are generally low (often <10%; e.g., Remage-Healey et al., 2008), and this affects the minimum time windows required for dialysate collection (often 5-30 min) in order to have a detectable amount of steroid. Steroid recoveries across the dialysis membrane in vivo are very difficult to estimate, and thus one cannot calculate absolute steroid levels in the extracellular fluid. Lastly, microdialysis probe insertion causes injury to the brain, which may induce neurosteroid synthesis (Garcia-Segura et al., 2001; Saldanha et al., 2009). This must be considered when interpreting the results.

Many studies have investigated steroids produced by brain tissue *in vitro*. These studies can examine steroid synthesis from radiolabeled precursors, as well as measure endogenous steroid levels in the conditioned medium and cultured tissue (organotypic brain slices, minces, or dissociated cells; Gomez-Sanchez et al., 1997; Holloway and Clayton, 2001; Hojo et al., 2004; Kretz et al., 2004; Munetsuna et al., 2009; Higo et al., 2011). These *in vitro* studies are very useful for investigating the regulation and functions of neurosteroids. However, the levels of steroids produced *in vitro* cannot directly reveal the levels of steroids in the brain *in vivo* (Schlinger et al., 1994; Saldanha et al., 2009). Moreover, creating brain slices or dissociating cells causes tissue injury and thus might affect steroid production.

EUTHANASIA

To directly measure steroid levels in various brain regions of animals, the brain must be collected after euthanasia of the subjects. Importantly, the method of euthanasia can affect steroid concentrations in the brain. The appropriate method of euthanasia should minimize stress and discomfort of the subjects as well as minimize changes in brain steroid levels. These considerations will vary depending on the nature of the experiment and the study species. A further consideration is whether blood should be collected and, if so, whether blood should be collected immediately before or after euthanasia and how and from where blood should be collected (Newman et al., 2008b).

Euthanasia is commonly performed by an overdose of an anesthetic. This method allows the animal to be subsequently perfused with saline (to remove blood from the brain; see below). However, inducing a loss of consciousness via anesthetic requires an alteration of neural activity (MacLusky, 2009), which might affect brain steroid levels. The effects of anesthesia on brain steroids can be specific to the anesthetic agent used. In rabbits, isoflurane or halothane gas anesthesia may rapidly increase plasma corticosterone and cortisol levels (González-Gil et al., 2006). However, in rats, carbon dioxide anesthesia for up to 2 min does not affect plasma corticosterone levels (Hackbarth et al., 2000). To our knowledge, no studies have examined the acute effects (within 2-3 min) of inhalant anesthesia on brain steroid measurements. Effects of injectable anesthetics on brain steroids also appear to be specific to the anesthetic used. For example, brain progestin levels are increased by injection of chloral hydrate or urethane, but brain progestin levels are decreased by pentobarbital, ketamine, or clonazepam (Korneyev et al., 1993a). Dramatic changes occur within 30 min, and in some cases within 10 min (Korneyev et al., 1993a). Again, however, we know of no studies examining the acute effects of injectable anesthetics on brain steroid measurements. Thus, when necessary and appropriate, anesthesia should be performed as rapidly as possible and with a method that minimizes changes in steroid levels. The stressors of handling and transfer into a novel cage for anesthesia may themselves cause rapid steroid changes. These stressors can be minimized by regular handling in the days prior to anesthesia or by administration of anesthesia in the home cage (e.g., Reburn and Wynne-Edwards, 2000).

Physical euthanasia without anesthesia does not depend on chemical alteration of nervous system activity (Karmarkar et al., 2010), and examples include decapitation or cervical dislocation without prior anesthesia (American Veterinary Medical Association, 2007). Loss of consciousness following decapitation is rapid (within seconds) and humane (Holson, 1992; Rijn et al., 2011), although some brain electrical activity can be recorded for up to 1 min (Rijn et al., 2011), which could potentially alter steroid levels. Rapid cooling of brain tissue might prevent such neurochemical changes, if present.

Euthanasia can also be performed by head-focused microwave irradiation, which rapidly heats the brain to cause extremely rapid loss of consciousness while maintaining physical integrity (Murphy, 2010). Heat inactivation stops enzymatic activity, thus rapidly halting lipid metabolism (Murphy, 2010) and changes in protein phosphorylation (O'Callaghan and Sriram, 2004), which otherwise continue after physical euthanasia. Focused microwave

irradiation is thus potentially useful for preventing continued steroid metabolism and has been used in a few studies of brain steroid levels (e.g., Korneyev et al., 1993a,b; Barbaccia et al., 1996; Concas et al., 2000).

To our knowledge, no studies have directly compared different methods of euthanasia and their effects on brain steroid levels. Such data would be very useful for researchers selecting the best method of euthanasia for an experiment.

PERFUSION WITH PHYSIOLOGICAL SALINE

In some studies, it may be important to remove blood contamination from the brain. This can be achieved via transcardial perfusion with physiological saline (following anesthesia). For example, saline perfusion is often performed prior to tissue fixation for immunohistochemistry. Also, removal of blood contamination might give a more accurate estimate of steroid concentrations in the brain tissue (i.e., without steroids in the blood; Taves et al., 2010). This issue has been experimentally examined in rats, mice, and songbirds. In neonatal rats and adult mice, saline perfusion has no effects on brain 17β-estradiol and corticosterone levels, respectively (Amateau et al., 2004; Little et al., 2008). However, in adult zebra finches, saline perfusion increases brain corticosterone levels, decreases brain DHEA levels, and increases brain 17β-estradiol levels in a region-specific manner, possibly due to rapid effects of ischemia (Taves et al., 2010). Such steroid changes might be minimized by perfusion with oxygenated artificial CSF. Thus, when necessary, perfusion should be performed as rapidly as possible. Furthermore, because saline perfusion does not cause a global decrease in brain steroid levels, blood contamination (\sim 1% of brain, v/w) does not appear to significantly affect brain steroid measurements (Taves et al., 2010). Prior to an experiment, validations should be performed to ensure that saline perfusion does not affect brain steroid levels.

DISSECTION

For measurement of brain steroid concentrations, nearly all studies have used gross dissection of tissue blocks that contain brain regions of interest (e.g., Butte et al., 1972; Caruso et al., 2008; Croft et al., 2008; Newman and Soma, 2009; Brummelte et al., 2010). To do this, the brain can be briefly cooled on wet ice, and then a scalpel blade is used to cut blocks of tissue based on their position relative to major neuroanatomical landmarks (e.g., midline, ventricles). Large areas, such as the cerebral cortex or hippocampus, can be subdivided as appropriate. Typically, tissue blocks are then frozen on dry ice and stored at -80° C. Such tissue blocks are relatively large, which increases the likelihood that steroids will be detectable in the sample and might also allow a greater number of steroids to be measured in a single sample. The major disadvantage of gross dissection is the loss of neuroanatomical specificity. Additionally, gross dissection by hand could increase the variability in the size of tissue samples across subjects.

Microdissection of specific brain regions allows for a higher degree of spatial specificity of brain steroid measurements (Magariños et al., 1989). Recently, microdissection of brain nuclei using the Palkovits punch technique (Palkovits, 1973; de Kloet, 2006) has been used for measurement of brain steroid concentrations (Charlier et al., 2010, 2011). Here, the whole brain is immediately

frozen on powdered dry ice and stored at -80° C. The brain is then covered in an embedding matrix intended for sectioning at a relatively warm temperature (e.g., Tissue-Tek O.C.T.), and brain sections (200–500 μ m) are cut on a cryostat at -12 to -10° C. The sections should be kept frozen at all times. Locations of specific brain nuclei can be determined using major neuroanatomical landmarks and a brain atlas. Sections containing nuclei of interest are mounted onto glass microscope slides, and cannulae (kept at -10° C or colder) are used to punch out tissue from the sections. The sections can then be stained with a Nissl stain to verify the locations of the punches (Charlier et al., 2010).

Microdissection has several advantages over gross dissection. Since the whole brain is frozen immediately after removal from the cranium, the amount of postmortem steroid metabolism is reduced. Additionally, microdissection allows for more precise targeting of specific brain nuclei. Finally, tissue samples are of a uniform size, which reduces between-subject variability in the amount of brain tissue per sample. A disadvantage of microdissection is that the tissue samples can be small (often <1 mg), and thus ultrasensitive assays are necessary to measure steroids in such small samples (Charlier et al., 2010). If the assays are also very precise, then samples need not be measured in duplicate (i.e., can be measured as singletons), which can be useful when the amount of sample is limited (Charlier et al., 2011). Note also that intermediate levels of spatial specificity can be achieved using a "brain matrix" (e.g., Ted Pella, #15007; Soma et al., 2005) or a tissue chopper (e.g., Ted Pella, #10180), followed by additional dissection with a scalpel or cannula.

Regardless of the method used to collect brain tissue, the amount of steroid in each sample must be corrected for the amount of tissue (typically wet weight in grams; see below). With gross dissection, each sample must be weighed on a microbalance. With Palkovits punch microdissection, however, all samples are of known thickness and diameter, and thus are of similar weight (Charlier et al., 2011). For example, when using a 1-mm-diameter cannula (Fine Science Tools, #18035-01) on 300 µm song sparrow brain sections, we have empirically determined that one brain punch has a mean wet weight of 0.245 mg (and protein content of 0.02 mg), with very little variation.

STEROID EXTRACTION

TISSUE HOMOGENIZATION

Once brain tissue samples have been collected, steroids must be extracted before they can be quantified (Newman et al., 2008a). This is especially critical in tissues such as brain that have very high lipid content, since lipids can interfere with steroid measurement (Rash et al., 1980). Prior to steroid extraction, tissue samples should be homogenized. To reduce potential metabolism of steroids in the tissue, tissue homogenization can be performed in ice-cold water and followed immediately by addition of an organic solvent such as methanol. Homogenization can also be performed in a water/methanol solution. Homogenization can be performed with a microcentrifuge tube and fitted pestle (Charlier et al., 2010), an electric hand-held homogenizer (e.g., Fisher Scientific PowerGen 125, #3396001; Newman and Soma, 2009; Taves et al., 2010), or bead mill homogenizer (e.g., Omni Bead Ruptor 24, #19-101). We have found that each of these

homogenization techniques results in a similar mean steroid concentration, although the bead mill homogenizer increases sample throughput and appeared to produce less variability across samples in a pilot study (unpublished results). Bead mill homogenization also prevents cross-sample contamination. To further ensure disruption of cells, tissue can be homogenized in lysis buffer (Konkle and McCarthy, 2011), or homogenized samples can be placed in a sonicator.

STEROID EXTRACTION

Following tissue homogenization, steroids must be extracted to separate them from other components of the brain sample, especially other lipids. Liquid–liquid extraction with organic solvents (e.g., diethyl ether) has long been used to isolate steroids, and is still frequently used. However, with liquid–liquid extraction, steroid recoveries can be low or variable, depending on the sample type and steroids of interest (Appelblad and Irgum, 2002). Liquid–liquid extraction can also extract other substances (i.e., other lipids in the brain; Appelblad and Irgum, 2002). These problems can impair accurate measurement of steroids in brain tissue samples. For these reasons, liquid–liquid extraction is being replaced by solid phase extraction over time (Makin and Gower, 2010), or can be paired with solid phase extraction (below; Chao et al., 2011).

Solid phase extraction of brain steroids typically involves loading the sample (in an aqueous buffer) onto a primed and equilibrated sorbent (e.g., silica-bonded C₁₈; Telepchak et al., 2004). We have found C₁₈ columns with 500 mg packing material (e.g., Agilent Bond Elut LRC-C18 OH, #12113045) mounted on a glass vacuum manifold (UCT #VMF024GL) to be suitable. After column priming and sample loading are completed, the sample is thoroughly washed to remove interfering substances (Appelblad and Irgum, 2002). With the correct wash(es), interfering substances can be effectively removed, while maintaining high steroid recoveries (Newman et al., 2008a). The selection of an effective wash step is dependent on the analyte, study species, tissue type, and method for steroid quantification (e.g., immunoassay, LC-MS/MS). Work in our laboratory, which has focused on unconjugated steroids, has found that washing with 10 ml deionized water is sufficient for songbird brain samples, but washing with 10 ml 40% methanol is required for mouse and rat brain samples (Figure 1). This might be related to species differences in brain lipid content. Note that 40% methanol also elutes conjugated (e.g., sulfated, glucuronidated) steroids (Bélanger et al., 1990; Wang et al., 1997; Liere et al., 2000). More stringent washes (e.g., 40% methanol, or deionized water followed by hexanes) may also be required if extracted steroids are to be separated by high-performance liquid chromatography (HPLC). After washing, steroids are slowly eluted off the sorbent into borosilicate glass vials or tubes. Elution of unconjugated steroids may be performed with 85-100% methanol, ethyl acetate, or one followed by the other (Bélanger et al., 1990; Wang et al., 1997; Scott and Ellis, 2007; Wong et al., 2008). Eluting unconjugated steroids with 85-90% methanol might be useful because 100% methanol can also elute steroids conjugated to fatty acids (Bélanger et al., 1990; Wang et al., 1997; Liere et al., 2004). It is critical to use the appropriate grade of solvents for solid phase extraction (e.g., HPLC-grade, MS-grade). Finally, solid phase extraction can be automated with robotics

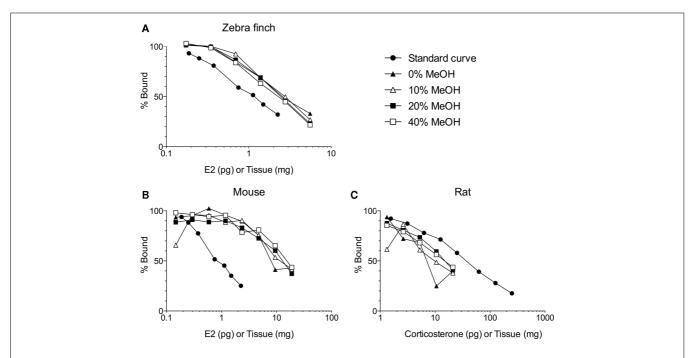


FIGURE 1 | Solid phase extraction of steroids from increasing amounts of brain tissue per assay tube. Samples were loaded onto C_{18} columns and washed with 10 ml of 0, 10, 20, or 40% methanol (MeOH) in water. Steroids were then eluted from the C_{18} columns with 5 ml of 90% MeOH. **(A)** The different washes have no effect on measurement of 17β-estradiol (E_2) in zebra finch brain (Taves and Soma, unpublished data). **(B)** The different washes do affect measurement of E_2 in mouse brain (Taves and Soma, unpublished data). **(C)** The different washes also affect measurement of corticosterone in rat brain (adapted with permission from Brummelte et al., 2010). The methods for all validations are similar (see Newman et al., 2008a). Briefly, brain tissue was homogenized in deionized water, and then MeOH was added. The

samples were mixed and then left overnight at 4°C. The samples were mixed again and centrifuged (3000 g, 10 min, 2°C). Supernatants (\leq 1 ml) were diluted with 10 ml deionized water and loaded onto C $_{18}$ columns (500 mg C $_{18}$, non-endcapped). Prior to sample loading, the C $_{18}$ columns were primed with 3 ml ethanol and equilibrated with 10 ml deionized water. After sample loading, columns were washed with 0, 10, 20, or 40% MeOH (10 ml) and steroids were eluted with 5 ml 90% MeOH. Eluates were dried in a vacuum centrifuge and resuspended in assay buffer. All reagents were HPLC-grade. E $_2$ was measured using an ultrasensitive and specific radioimmunoassay (Beckman, DSL-4800; Charlier et al., 2010), and corticosterone was also measured using a radioimmunoassay (MP Biomedicals, cat. 07120103, Washburn et al., 2002).

to increase sample throughput (e.g., Gilson #GX-274 ASPEC or Hamilton STARlet).

Solid phase extraction gives reliable recovery of very low steroid amounts from small samples. The combination of high steroid recovery and high precision make solid phase extraction an extremely useful tool (Telepchak et al., 2004). It is critical to routinely check the effectiveness of the extraction procedure. Note that some brands of C₁₈ columns can show lot-to-lot variation, and thus it is preferable to extract all the samples from one experiment with columns from the same lot. Furthermore, extraction performance should not be tested solely with radiolabeled steroids because this can overlook some problems. The use of only radiolabeled steroids ignores the potential co-elution of interfering substances (originating from the sample itself or from the C₁₈ column) and damage to the steroid analytes during processing (Scott and Ellis, 2007). Thus, it is important to routinely check extraction performance with unlabeled steroid standards ("quality controls") and water blanks.

After eluates have been collected, they can be dried at 40–60°C (Liere et al., 2000). We have found that drying at 50 or 60°C reduces drying time without affecting steroid measurement (**Table 1**; our unpublished results). Drying is typically done under a stream of high-purity nitrogen with a sample concentrator (e.g.,

Techne Dri-Block DB-3) or in a vacuum centrifuge (e.g., Thermo Scientific SPD111V SpeedVac). The dried steroid residue can then be resuspended for quantification, for example via immunoassay. We have found that a small amount of absolute ethanol improves steroid resuspension into an aqueous buffer (Newman et al., 2008a), but care must be taken because ethanol (or impurities therein) can affect some immunoassays (Ocvirk et al., 2009; Charlier et al., 2010). Alternatively, after solid phase extraction, steroids can be separated via chromatography prior to quantification (see below).

CHROMATOGRAPHIC SEPARATION

CELITE COLUMN CHROMATOGRAPHY

The C_{18} columns used for solid phase extraction are relatively short, and although they can separate conjugated steroids from unconjugated steroids (Liere et al., 2004), they have little ability to separate different unconjugated steroids from each other (e.g., separate testosterone from estradiol). In many cases, it is useful to separate the steroids present in a brain tissue sample prior to steroid measurement. Steroid separation allows several steroids to be measured (e.g., via immunoassays) without splitting the sample, which is useful when the tissue sample is small. Furthermore, with immunoassays, antibody cross-reactivity can be a

Table 1 | Effect of drying temperature on measured steroid values.

40°C (%)	50°C (%)	60°C (%)		
100	114	101		
100	108	113		
100	96	103		
	100	100 114 100 108		

Samples were dried in a preheated Thermo Scientific SPD111V SpeedVac. Samples dried at 40°C (the most common temperature used across many laboratories) were used as the baseline, and samples dried at 50 and 60°C are expressed as a percent of this baseline. Each value is the mean of three samples. Drying at 40, 50, and 60°C took approximately 3.0, 2.5, and 2.0 h, respectively. Taves and Soma, unpublished data.

concern (see below), and this concern can be alleviated by steroid separation prior to assay (e.g., Close et al., 2010).

Celite column chromatography uses columns or cartridges containing celite (highly porous, diatomaceous earth). Samples are loaded onto celite columns, and then different steroids are eluted off with different organic solvents. Celite chromatography has sufficient resolving power to separate multiple steroids and has been used in numerous studies of plasma and brain steroids (Wingfield and Farner, 1975; Corpéchot et al., 1981; von Schoultz et al., 1990; Wang et al., 1997; Frye et al., 2000; Dinger and Frye, 2004; Rosario et al., 2004; Pradhan et al., 2010b). Nonetheless, celite columns have less resolving power than HPLC or gas chromatography.

HIGH-PERFORMANCE LIQUID CHROMATOGRAPHY

High-performance liquid chromatography is often used to separate steroids in a sample. Reversed-phase HPLC using a C_{18} HPLC column (e.g., Waters SymmetryShield, 4.6 mm × 250 mm, 5 μ m silica particles) is often used for such purposes, allowing multiple steroids to be collected and measured, without the need to split the sample (**Figure 2**). Furthermore, for immunoassays, HPLC separation removes potentially cross-reactive steroids, resulting in greater confidence in immunoassay measurements. HPLC often requires minimal sample preparation, as steroids need not be derivatized beforehand. This allows for fraction collection and quantification by immunoassay (Mensah-Nyagan et al., 1996; Munetsuna et al., 2009), more extensive purification and analysis (Mensah-Nyagan et al., 1996; Liere et al., 2000, 2004), or subsequent analysis via tandem mass spectrometry (Liu et al., 2003; Krone et al., 2010).

GAS CHROMATOGRAPHY

Steroid separation can also be done via gas chromatography, prior to mass spectrometry (Vallée et al., 2000; Alomary et al., 2001). Gas chromatography requires some sample preparation, as steroids are generally derivatized prior to GC/MS analysis. While derivatization can greatly increase sensitivity and aid in determination of steroid structure, it can be a time-consuming process and can convert conjugated steroids into unconjugated steroids (Soldin and Soldin, 2009) or even convert cholesterol into steroids (Schumacher et al., 2008; Liere et al., 2009). Different conjugated steroids (e.g., DHEA-sulfate and DHEA-glucuronide) may also be converted into the same form, resulting in conjugate misidentification (Liu et al., 2003). Additionally, the derivatization process is challenging to automate, which reduces sample throughput (Krone

et al., 2010). Nonetheless, GC/MS is well suited for the identification of novel steroidal compounds (Krone et al., 2010). Further, the National Institute of Standards and Technology (NIST) GC/MS database search capability is a powerful tool to aid in the discovery and analysis of novel steroids.

STEROID QUANTIFICATION

IMMUNOASSAYS

Immunoassays include both radioimmunoassays (RIA) and enzyme-linked immunosorbent assays (ELISA, also known as, EIA). Immunoassays are still the most common way to measure steroids, because of their rapidity, simplicity, sensitivity, and cost-effectiveness. Furthermore, many RIA and ELISA kits for steroids are commercially available, contributing to the popularity of immunoassays. Many commercially available RIA kits can be modified to greatly increase their sensitivity. For example, we have modified a double-antibody 125 I RIA kit to detect sub-picogram amounts of 17β -estradiol (Charlier et al., 2010). This modified assay is also accurate, specific, and precise. Samples are usually measured in duplicate, requiring the sample to be split in half, but the high precision of this estradiol assay allows samples to be run as singletons, which is useful when the sample is limited (Charlier et al., 2011).

There are some disadvantages associated with the use of immunoassays. One limitation is antibody cross-reactivity, particularly with polyclonal antibodies. Antibodies exhibit widely varying degrees of specificity for their antigen and may bind molecules other than the target of interest, such as closely related steroids (Chard, 1995). Several measures can alleviate concerns about antibody cross-reactivity: selection of a specific antibody, checking closely related steroids for antibody binding, and chromatographic separation of steroids prior to immunoassay.

Another potential problem is that different matrices can affect antibody binding. Most commercially available steroid immunoassay kits are designed to measure steroids in unextracted plasma, serum, or saliva samples, and thus the standards in the standard curve contain the steroid in an appropriate matrix (i.e., similar to serum or saliva). However, when steroids in brain samples are extracted and resuspended in an aqueous buffer that does not contain serum or saliva components (e.g., phosphate-buffered saline with gelatin, PBSG), the antibody can show increased binding to the analyte (possibly due to reduced matrix interference). Thus, extracted steroids resuspended in PBSG should be compared to a standard curve that is also diluted in PBSG. These "matrix effects" can affect steroid measurements, especially at low steroid concentrations.

MASS SPECTROMETRY

Mass spectrometry, following either gas or liquid chromatography, is highly specific and is currently the "gold standard" for definitive identification of analyte molecules (Alomary et al., 2001; Soldin and Soldin, 2009; Makin and Gower, 2010). Mass spectrometry also allows for simultaneous measurement of many steroids in a single sample, or "steroid profiling" (Liu et al., 2003; Ebner et al., 2006; Meffre et al., 2007; Soldin and Soldin, 2009; Koren et al., submitted). This can now be used to measure both unconjugated steroids and intact steroid sulfates and glucuronides (Jäntti et al.,

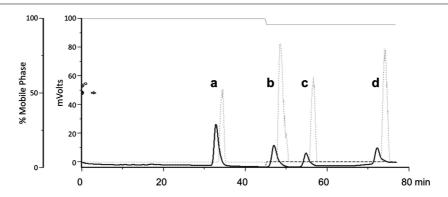


FIGURE 2 | High-performance liquid chromatography separation of (a) cortisol, (b) corticosterone, (c) 11-deoxycortisol, and (d)

11-deoxycorticosterone. To separate these glucocorticoids, we used reversed-phase HPLC (Gilson 322) with a Waters SymmetryShield C_{18} column (4.6 mm \times 250 mm, 5 μ m silica particles). The mobile phase consisted of 50% HPLC-grade methanol in HPLC-grade water with 0.01% glacial acetic acid (solvent A; indicated by the solid gray line at top). At 45 min, 5% isopropanol (solvent B; dashed gray line at the bottom) was added in order to elute the

final peaks more rapidly. The solid black line indicates radioinert steroids measured with a UV detector, and the dotted gray line indicates 3 H-labeled steroids measured with a radioflow detector (as described in Mensah-Nyagan et al., 2008; Pradhan et al., 2008, 2010a). Detection of radiolabeled steroids occurs 1.6 min later than radioinert steroids because of the extra length of tubing between the UV and radioflow detectors. A fraction collector was then used to collect glucocorticoids for quantification via enzyme immunoassay (EIA) or radioimmunoassay (Taves et al., unpublished results).

2010). These characteristics have contributed to the growing use of mass spectrometry (and tandem mass spectrometry) in steroid analysis (Mensah-Nyagan et al., 1996; Griffiths et al., 2001; Liere et al., 2004, 2009; Matsunaga et al., 2004; Wang et al., 2007; Caruso et al., 2008; Krone et al., 2010).

The sensitivity of mass spectrometry has greatly improved. Steroids are typically derivatized prior to GC/MS to increase measurement sensitivity, but this has some disadvantages (see above). Steroids need not be derivatized prior to LC-MS/MS, although this can be done to increase sensitivity and specificity (Higashi and Shimada, 2004; Higashi et al., 2005; Higashi, 2006; Hojo et al., 2009). Since the sensitivity of LC-MS/MS varies for different steroids, derivatization can be used to aid detection of poorly ionized steroids when they are present in low quantities. A variety of derivatization methods can be used, depending on the analyte(s) of interest (Santa, 2011). The method of ion production can also affect sensitivity (Wang et al., 2007). For some steroids, atmospheric pressure photo-ionization might yield greater sensitivity than atmospheric pressure chemical ionization or electrospray ionization (Robb et al., 2000; Guo et al., 2006). Nonetheless, when the amount of brain tissue is extremely limited (e.g., punch samples or samples from neonatal rats) the limits of quantification of mass spectrometry may not be sufficient (Konkle and McCarthy,

Thus, mass spectrometry provides definitive steroid identification, simultaneous multi-analyte quantification and good sensitivity. However, high equipment costs and operating costs and the need for specific technical expertise are obstacles to using mass spectrometry for steroid analysis (Makin and Gower, 2010; Stanczyk and Clarke, 2010).

INTERPRETATION

COMPARING STEROID LEVELS IN THE BRAIN AND CIRCULATION

Many studies directly compare steroid levels in brain tissue with steroid levels in the general circulation. In subjects that have been adrenalectomized and/or gonadectomized for a long time, the presence of a steroid in the brain, but not in the circulation, is strong evidence for local steroid synthesis (Corpéchot et al., 1981; Concas et al., 2000; Gomez-Sanchez et al., 2005; Croft et al., 2008; Higo et al., 2011). Furthermore, in gonad- and adrenal-intact subjects, if tissue steroid levels are higher than circulating steroid levels, then these data suggest that steroids are locally synthesized, locally sequestered by tissue binding sites, or locally released from steroid binding proteins in the blood. These alternative possibilities can be tested, for example, by measuring steroidogenic enzymes and/or steroid binding sites (Taves et al., 2011). Some representative measurements of endogenous steroid levels in intact adult male rats are given in **Table 2**.

When comparing steroid levels in the brain and circulation, a critical point is that the use of plasma samples significantly overestimates steroid concentrations in the blood. Steroid concentrations in plasma are roughly twice as high as steroid concentrations in whole blood (Taves et al., 2010). Thus, for a more representative measurement of actual circulating steroid levels, it is informative to measure steroids in whole blood samples as well as plasma samples (**Figure 3**).

Another issue is that most studies measure total steroid levels (free + bound) in brain and plasma samples. In the case of corticosterone and cortisol, over 90% of these glucocorticoids in plasma are bound with high affinity to corticosteroid-binding globulin, and are thus unable to readily enter tissues (Hammond, 1990; Breuner and Orchinik, 2002; Charlier et al., 2009). Since only free (unbound) steroids are available to enter tissues, plasma free steroid levels are also informative for comparison to tissue steroid levels (as in Little et al., 2008; **Figure 4**).

If local steroid synthesis occurs at low levels or only in specific cells or subcellular compartments (e.g., mitochondria, presynaptic terminals), then the average tissue steroid concentration may be similar to or lower than the circulating steroid concentration (Newman and Soma, 2011). For all of the above reasons, when

Table 2 | Selected measurements of endogenous steroid levels in adult male rat plasma or serum (ng/ml) and brain tissue (ng/g).

Steroid	Plasma/serum	Cortex	Hippocampus	Hypothalamus	Cerebellum	Whole brain
Descriptions	- 1 00	0.4–7.0	4.0	11–12	0.4–19	17.40
Pregnenolone	n.d.–0.9	0.4-7.0	4.8	11-12	0.4-19	1.7–4.9
Progesterone	0.5–2.2	1.5–7.0	4.6	0.9	0.7–5.3	0.4–0.7
DHEA	n.d.–0.1	0.2	0.1	1.5	0.2	0.2-0.3
Testosterone	1.2-4.2	0.1–2.6	4.9	14	2.3-2.4	1.4-2.2
17β-Estradiol	n.d0.004	0.1	2.3	n.a.	0.02	n.d.
5α-DHT	n.d0.2	n.d0.2	1.9	2.0	0.1	n.d.
Corticosterone	20–54	7.5	44	43	20	4.6–50

All brain steroid measurements were obtained after solid phase extraction. In some cases, steroid extracts were further purified using high-performance liquid chromatography, gas chromatography, or thin-layer chromatography. Steroids were quantified using radioimmunoassay or mass spectrometry. DHEA, dehydroepiandrosterone; 5\(\textit{5a}\)-DHT, 5\(\textit{a}\)-dihydrotestosterone; n.d., non-detectable; n.a., not assessed. Values compiled from Butte et al. (1972); Robel et al. (1973); Lanthier and Patwardhan (1987); Purdy et al. (1991); Korneyev et al. (1993b); Wang et al. (1997); Vallée et al. (2000); Ebner et al. (2006); Meffre et al. (2007); Caruso et al. (2008); Hojo et al. (2009): Pesaresi et al. (2010).

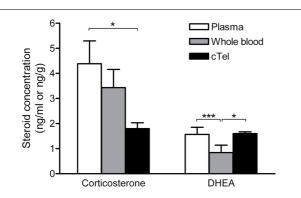


FIGURE 3 | Corticosterone and DHEA levels in zebra finch plasma, whole blood, and brain (cTel, caudal telencephalon). For corticosterone, brain levels are significantly lower than plasma levels but not significantly lower than whole blood levels. For DHEA, brain and plasma levels are similar, but brain levels are significantly higher than whole blood levels. Thus, the use of plasma samples (which overestimate circulating steroid levels) or whole blood samples can lead to very different interpretations when comparing tissue steroid levels against circulating steroid levels. *p < 0.05, ***p < 0.001. Reprinted with permission from Taves et al. (2010).

total steroid levels in brain tissue are similar to or even lower than total steroid levels in plasma, one cannot exclude the possibility of local steroid synthesis on the basis of this observation alone. Other forms of evidence (e.g., steroidogenic enzyme expression) are useful for interpreting such a pattern of results.

To facilitate comparison of brain and circulating steroid levels, it is extremely useful to present brain steroid concentrations and circulating steroid concentrations in comparable units (e.g., ng/g wet weight for brain, ng/ml for plasma or whole blood). Note that 1 ml of plasma or blood weighs very close to 1 g (Schmidt et al., 2008; Taves et al., 2010). Some papers use different units for brain steroid concentrations and plasma steroid concentrations (e.g., ng/mg protein for brain, nM for plasma). This practice complicates the comparison of brain and plasma steroid concentrations.

COMPARING STEROID LEVELS IN DIFFERENT BRAIN REGIONS

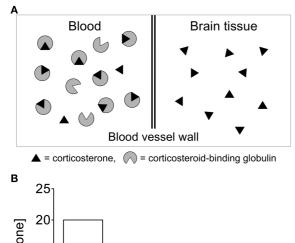
Within an individual brain, a comparison of steroid levels in different regions can identify which regions are most active as steroid producers or targets (von Schoultz et al., 1990; Toran-Allerand et al., 2005; Little et al., 2008; Newman and Soma, 2009; Charlier et al., 2010). This is important in identifying region-specific effects of experimental treatment. For example, if an experimental treatment elevates steroid levels in a specific brain region, but not in other brain regions or circulating blood, this indicates local steroid production (or accumulation) in that specific region (e.g., Little et al., 2008). Comparing steroid levels across brain regions is somewhat more straightforward than comparing steroid levels between brain and blood.

EFFECTS OF PERIPHERAL STEROIDS ON BRAIN STEROID LEVELS

Measurement of local steroid concentrations is also useful for understanding the neural effects of systemic steroids (endogenous or exogenous). It is generally assumed that increases in systemic steroid levels result in concurrent increases in steroid levels in target tissues, but this is not necessarily the case. For example, in rats, a rise in plasma corticosterone is followed by a rise in brain corticosterone (as measured by microdialysis), but surprisingly only after a 20-min delay (Droste et al., 2008). Systemically administered steroids may also be metabolized before reaching the brain or within the brain itself (Edinger and Frye, 2004), and some steroids are preferentially excluded from the brain via active transport by P-glycoprotein in the blood-brain barrier (Karssen et al., 2001). By measuring steroid levels in target tissues (such as brain), researchers can test whether a systemic steroid treatment results in locally elevated steroid levels, and whether local steroid levels are within the physiological range. Furthermore, as some protein-bound circulating steroids have minimal access to the brain (Pardridge and Mietus, 1979), systemic steroid treatments that result in supraphysiological levels in the blood may produce only moderately elevated steroid levels in the brain (e.g., Amateau et al., 2004). Finally, note that systemically administered steroids can regulate the synthesis of neurosteroids (e.g., Soma et al., 2004).

INADVERTENT INDUCTION OF STEROIDOGENIC ENZYMES IN THE BRAIN

Injuries to brain tissue can have rapid and enduring effects on steroidogenic enzyme activities around the damaged brain tissue.



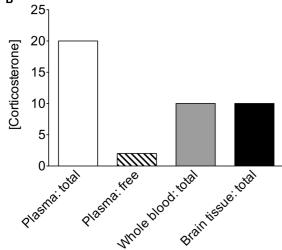


FIGURE 4 | Theoretical comparison of circulating and tissue steroid concentrations. (A) If the same amounts of steroid are present in equal volumes of blood and brain, different interpretations can be made, depending on how steroids are quantified in blood. In the circulation, the vast majority of corticosterone is bound to corticosteroid-binding globulin (CBG; Hammond, 1990). CBG-bound steroids have very limited access to tissue and limited ability to cross the blood-brain barrier. (B) Corticosterone concentrations in whole blood are approximately 50% of those in plasma (Taves et al., 2010), and within plasma, the vast majority of corticosterone is bound to CBG. In the above theoretical scenario, if whole blood corticosterone is measured, then circulating levels appear identical to tissue levels. If plasma total corticosterone is measured, then circulating levels appear much higher than tissue levels. If plasma free corticosterone is measured, then circulating levels appear much lower than tissue levels. Thus, the use of whole blood steroid measurements or plasma free steroid measurements is highly informative in addition to plasma total steroid measurements

For example, aromatase is expressed constitutively in avian and mammalian neurons (Balthazart et al., 1993; Naftolin et al., 1996; Saldanha et al., 2000), but neuropil perturbation, such as by excitotoxicity (Azcoitia et al., 2001), penetrating injury (Garcia-Segura et al., 1999; Peterson et al., 2001), or anoxia/ischemia (Garcia-Segura et al., 2001), induces additional aromatase expression in astroglia around the site of primary damage (Saldanha et al., 2010). This induction of aromatase occurs in hours and lasts weeks (Wynne et al., 2008) or months post-injury (Garcia-Segura et al., 2001). Importantly, this induction occurs in a cell type that only expresses aromatase *in vitro* or following

damage to the brain (Schlinger et al., 1994; Zwain and Yen, 1999; Saldanha et al., 2009). Thus, when experimental techniques that cause physical damage of neural tissue are used, measured steroid levels likely reflect a combination of constitutive and induced estrogen levels. Such techniques could include brain slicing or dissociation prior to tissue culture, microdialysis probe or cannula insertion, saline perfusion, or a delay between euthanasia and brain freezing. Increases in brain estradiol levels occur within minutes of insult (Saleh et al., 2004; Taves et al., 2010). Other steroids might also be altered during routine tissue collection procedures.

CONCLUSION AND FUTURE DIRECTIONS

Numerous studies have measured steroids in plasma or serum, and many studies have examined steroidogenic enzyme expression in the brain. However, far fewer studies have measured steroids in the brain. For example, many more studies have measured aromatase than estradiol levels in the brain. However, measurement of tissue steroid concentrations is critical to understanding the numerous roles and regulation of steroid signaling molecules in the brain. Such measurements involve tissue collection, steroid extraction, and steroid separation and quantification. At each of these steps, there are various methodological approaches that may be used, and the selection of a particular approach necessitates weighing the costs and benefits of each, with respect to the goals of the particular experiment. If the results obtained with different approaches converge to give the same conclusion, then this is strong evidence for a given phenomenon. Thus, the use of various methods within a laboratory or across different laboratories can be useful

Improvements in steroid extraction and detection are steadily increasing our ability to measure extremely small quantities of steroids (<0.5 pg of steroid in a sample). Improved measurement sensitivity allows for techniques that measure steroids in samples that are of minimal size (e.g., brain punch samples, smaller microdialysate samples, CSF samples from human infants, tumor biopsies), which can improve the spatial and temporal resolution of brain steroid measurements. Laser capture microdissection gives excellent spatial specificity (de Kloet, 2006), and improved assay sensitivity might permit use of this technique for steroid quantification in brain tissue. Similarly, *in vivo* microdialysis provides temporal information about brain steroid changes, and improved assay sensitivity will permit the use of shorter dialysate collection windows, providing greater temporal resolution.

In the future, other methods of measuring steroids in the brain could be useful. For example, immunohistochemistry might allow for even greater spatial information about steroids in the brain, particularly at synapses. A few studies have quantified steroids via immunohistochemistry using specific antibodies in fixed neural and non-neural tissues (Dornhorst and Gann, 1978; Saalmann et al., 2007; Tokuda et al., 2011). It is critical that the necessary controls be run (e.g., pre-absorption controls, tissues lacking the steroid of interest) to convincingly demonstrate the specificity of antibody binding in sample tissues.

In addition to quantifying well-known steroids, future studies will undoubtedly discover novel steroids, steroid conjugates,

and steroid metabolites. GC/MS is especially suited for such exploratory analyses (Krone et al., 2010). The need for identification of novel steroids and steroid metabolites is well illustrated by the case of brain estradiol, whose downstream metabolism remains unclear (Cornil, 2009; Charlier et al., 2011). Note that local steroid metabolism is likely to affect local steroid levels (Jäntti et al., 2010; Charlier et al., 2011). Such studies, along with improvements in steroid profiling and measurement sensitivity, promise to provide a much more detailed understanding of regional and cellular differences in steroid synthesis, action, and metabolism in the brain.

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