

COMMENTARY

Good news for CB₁ receptors: endogenous agonists are in the right place

M Maccarrone^{1,2}

¹Department of Biomedical Sciences, University of Teramo, Teramo, Italy and ²European Center for Brain Research (CERC)/IRCCS S Lucia Foundation, Rome, Italy

Endocannabinoids are endogenous ligands of brain-type (CB₁) and spleen-type (CB₂) cannabinoid receptors. *N*-Arachidonylethanolamine (anandamide, AEA) and 2-arachidonoylglycerol (2-AG) are prototype members of the fatty acid amides and the monoacylglycerols, two groups of endocannabinoids. Unlike CB₁, CB₂ receptors do not reside within 'caveolae', specialized membrane microdomains that are well-known modulators of the activity of a number of G protein-coupled receptors. In this issue of the *British Journal of Pharmacology*, Rimmerman and coworkers demonstrate that 2-AG is entirely localized in the caveolae of dorsal root ganglion cells, where also part of AEA (~30%) can be detected. However, most of AEA (~70%) was detected in non-caveolae fractions, that is where CB₂ receptors are localized. The different interaction of AEA and 2-AG with membrane microdomains might have significant implications for endocannabinoid-dependent autocrine and/or retrograde-paracrine signalling pathways. It also raises an important question about the structural determinants responsible for a different localization of two apparently similar endocannabinoids within lipid bilayers.

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Abbreviations: AEA, *N*-arachidonylethanolamine (anandamide); 2-AG, 2-arachidonoylglycerol; CB₁, brain-type cannabinoid receptor; CB₂, spleen-type cannabinoid receptor; LRs, lipid rafts; QSAR, quantitative structure–activity relationship

Endocannabinoids are lipid signalling molecules that modulate several physiological processes. They are endogenous ligands of brain-type cannabinoid receptors (CB₁) and spleen-type cannabinoid receptors (CB₂), two G protein-coupled receptors that also bind Δ^9 -tetrahydrocannabinol, the psychoactive component of *Cannabis sativa* (Howlett *et al.*, 2002). *N*-arachidonylethanolamine (anandamide, AEA) and 2-arachidonoylglycerol (2-AG) are prototype members of two groups of endocannabinoids, the fatty acid amides and the monoacylglycerols respectively (Piomelli, 2003; Di Marzo and Petrosino, 2007).

The dependence of endocannabinoid signalling on 'lipid rafts' (LRs) is an emerging concept (Barnett-Norris *et al.*, 2005; Dainese *et al.*, 2007). LRs are specialized membrane microdomains that are enriched in cholesterol, sphingolipids and arachidonic acid and that have a tightly packed state (Hanzal-Bayer and Hancock, 2007). LRs are well-known modulators of the activity of a number of G protein-coupled receptors, for which a raft domain provides a more organized

platform for the proper assembly of signalling complexes, also preventing cross-talks between different pathways. CB₁ receptors have been shown to reside within LRs (Bari *et al.*, 2005), and consistently they co-localize with LR markers (Sarnataro *et al.*, 2006). More recently, CB₁ receptors have been shown to co-localize with caveolin-1, a marker of 'caveolae' (Bari *et al.*, 2007). These are considered a subclass of LRs represented by non-clathrin-coated and flask-shaped invaginations (diameter of ~60–80 nm) in the plasma membrane. Unlike CB₁, CB₂ receptors do not reside within LRs or caveolae (Bari *et al.*, 2006), as also shown by Rimmerman *et al.* (2008) in this issue of the *British Journal of Pharmacology*. These authors provide further information that seems to add a new player in the arena of the modulation of endocannabinoid signalling. In fact, they demonstrate by liquid chromatography/tandem mass spectrometry that 2-AG is concentrated in the caveolae of dorsal root ganglion cells where it co-localizes with components of the diacylglycerol pathway responsible for 2-AG production. Instead, AEA was detected in LRs and non-LR fractions to comparable levels (Rimmerman *et al.*, 2008). Therefore, much similar cannabinoid receptors (CB₁ versus CB₂), their endogenous ligands (AEA versus 2-AG) show a different interaction with LRs, an observation that might have significant implications for endocannabinoid-dependent

Correspondence: Professor M Maccarrone, Department of Biomedical Sciences, University of Teramo, Piazza Aldo Moro 45, Teramo 64100, Italy.
E-mail: mmaccarrone@unite.it
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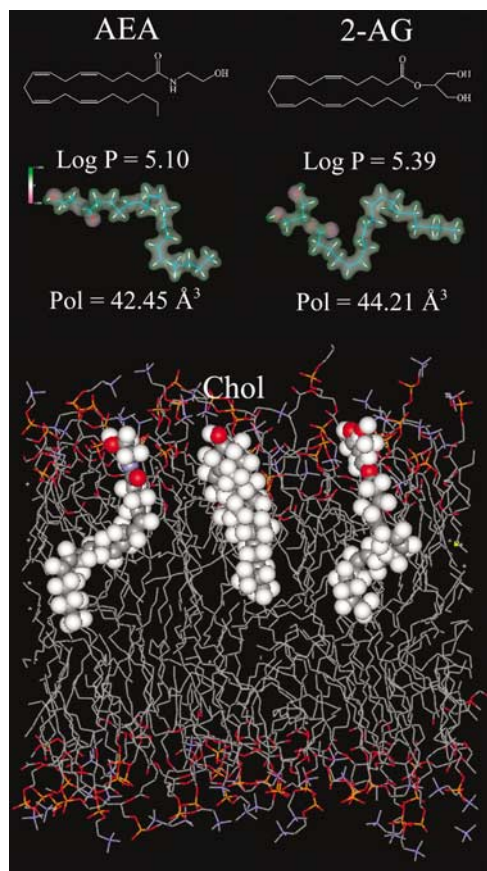


Figure 1 AEA versus 2-AG. The chemical structures, lipophilicity ($\log P$), electrostatic potential (violet -1 ; green $+1$) and polarizability (Pol) values of AEA and 2-AG are shown in the upper panel. Both $\log P$ and Pol values were calculated by means of the HyperChemTM 6.03 molecular modelling system (Hypercube, Inc., Gainesville, FL, USA), as reported (Dainese *et al.*, 2005). A schematic representation of (from left to right) AEA, cholesterol (Chol) and 2-AG embedded within a dipalmitoylphosphatidylcholine bilayer is shown in the lower panel. Here, atoms in the space-filling models were coloured with the following codes: oxygen in red, nitrogen in violet, carbon in grey and hydrogen in light grey. The figure was kindly provided by Dr. Enrico Dainese (University of Teramo, Teramo, Italy).

autocrine and/or retrograde-paracrine signalling. On one hand, the findings presented by Rimmerman *et al.* (2008) demonstrate that the CB₁ receptor agonists 2-AG and, to a lesser extent, AEA are present where they should be in order to activate their target: next to it, so that they can easily reach the binding site by lateral diffusion. On the other, these new data call for a reconsideration of the general concept that 2-AG is the only true agonist of CB₂ receptors whereas AEA (a weak and partial ligand for this receptor) does not have any physiological relevance (Sugiura *et al.*, 2000). In fact, AEA might be localized more likely than 2-AG next to CB₂ receptors in dorsal root ganglion cells and possibly in other cell types.

The paper by Rimmerman *et al.* (2008) raises another important question: 'what are the structural determinants responsible for the different localization of AEA and 2-AG within lipid bilayers?' Like many other bioactive lipophilic molecules, endocannabinoids partition into membranes

where they assume a thermodynamically favourable orientation and location. Quantitative structure–activity relationship (QSAR) studies have demonstrated that AEA and 2-AG share similar values of molecular descriptors like lipophilicity (expressed as logarithm of the octanol–water partition coefficient, $\log P$), distribution of electrostatic potential and polarizability (Dainese *et al.*, 2005). QSAR data were in accordance with the high flexibility of AEA observed by molecular dynamics simulations, which also showed that AEA embedded within the lipid bilayer tends to adopt a more extended conformation (Barnett-Norris *et al.*, 2002). On the basis of the similarities of the molecular descriptors derived from QSAR analysis, it seems that 2-AG is embedded in the lipid bilayer in the same manner as AEA (Figure 1). Therefore, it is rather unexpected that these two endocannabinoids partition quite differently in raft and non-raft fractions. Maybe cholesterol, that is known to bind with AEA (Biswas *et al.*, 2003), binds with 2-AG even better, presumably due to a better interaction with its acyl chain (Figure 1). As a consequence, cholesterol may favour the concentration of 2-AG within LRs. However, the study by Rimmerman *et al.* (2008) underlines the need for extensive investigations into the structural determinants that drive the interaction of apparently similar endocannabinoids with the surrounding lipid environment. Furthermore, the dependence of CB₁ receptors on LRs integrity makes it challenging from the therapeutic point of view to selectively target CB₁-dependent pathologies by means of LRs-oriented drugs. On a final note, the different interaction of AEA and 2-AG with raft and non-raft microdomains, along with the different localization of CB₁ and CB₂ receptors within these fractions, might represent a novel paradigm of ligand–receptor interactions whereby a third player comes into the game—the membrane lipids.

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