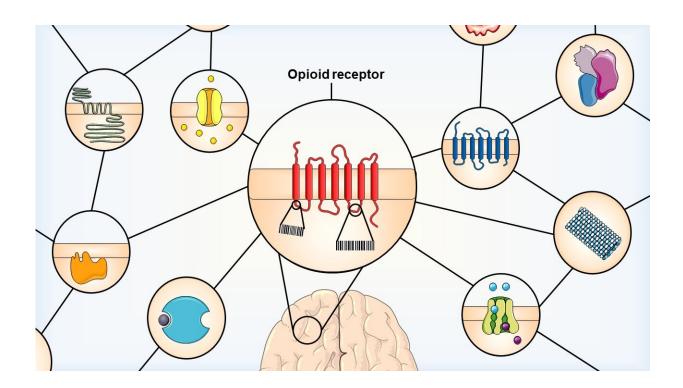


Study examines the differences between opioid receptor subtypes

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Each opioid receptor subtype displays specific sequences or "barcodes" recognized by regulatory proteins. This unique network of interacting proteins will dictate its localization and signaling, thus impacting its physiological functions such as induced analgesic effects. Credit: Jade Degrandmaison.

Opioid receptors are proteins in the central nervous system that are targeted by opioid drugs for pain relief. There are three subtypes of opioid receptors, MOP, DOP and KOP, in addition to the opioid-related



NOP receptor.

The most prescribed opioids, such as morphine, fentanyl and codeine, target μ opioid receptors (MOP). In recent years, however, many neuroscientists have also been investigating the therapeutic potential of drugs that target the other subtypes of opioid receptors.

Past research suggests that the δ -opioid receptor (DOP) could be a particularly promising therapeutic target for chronic pain management drugs. In fact, substances targeting DOP receptors appear to have fewer adverse effects than commonly used opioids, while still providing relief for pain associated with inflammation, diabetes, cancer and other health conditions.

While the molecular mechanisms governing the <u>trafficking</u> of MOP receptors are now well understood, there have so far been relatively few studies focusing on the trafficking of DOP receptors. Some recent papers, however, have highlighted a series of unique characteristics that could set DOP receptors apart from other subtypes of opioid receptors.

Researchers at Université de Sherbrooke in Canada have recently reviewed these past findings with the aim of painting a more exhaustive picture of the mechanisms governing the trafficking of DOP receptors. Their paper, published in the *Journal of Neuroscience Research*, summarizes what is currently known about the amino acid patterns involved in the regulation of different opioid receptors' trafficking.

"Over the past several years, we have been interested in the <u>molecular</u> <u>mechanisms</u> underlying the distinctive regulation of the δ -opioid receptor, a promising pharmacological target for the treatment of chronic pain as well as various mood/<u>psychiatric disorders</u>," Louis Gendron, one of the researchers who carried out the study, told *Medical Xpress*. "Although DOP shares high homology with the other opioid



receptor subtypes, this receptor displays very unique characteristics regarding its subcellular localization and intracellular trafficking."

The mechanisms underlying the intracellular trafficking of opioid receptors are known to depend on the receptors' interactions with regulatory proteins. Therefore, Gendron and his colleagues set out to delineate the similarities and dissimilarities between amino acid sequences and motifs (or "barcodes") that could explain the differences between DOP receptors and other opioid receptors.

"Given their distinctive characteristics, many studies have used DOP and MOP as model receptors for the study of trafficking mechanisms," Gendron explained. "We thus took advantage of these comparative studies to highlight and discuss the differences between both receptors."

Gendron and his colleagues also reviewed studies that investigated the trafficking of other G protein-coupled receptors (GPCRs), as the "barcodes" identified in these studies could also be relevant to the opioid receptor physiology. Although less is known about the regulation of their trafficking, the team also discussed the trafficking of kappa (KOP) and nociceptin (NOP) receptors.

"While writing our review, we kept in mind that our analysis and mapping of the known recognition and regulatory motifs ("barcodes") could help future investigations of these important receptors," Gendron said.

The literature review authored by this team of researchers could have important implications. Most notably, it identified a new series of potential "barcodes" that could be specific to each opioid receptor subtype, which could be further examined in future studies.

"Although further research will be needed to validate our predictions, we



think that our analysis could help guide future investigations," Gendron said.

In the future, the researchers' paper could prove to be a valuable resource for neuroscientists investigating the trafficking of <u>opioid</u> <u>receptors</u> and of GPCRs in general. In fact, other receptors in the central nervous system could be regulated by cellular mechanisms that resemble those outlined by Gendron and his colleagues.

"We are now investigating several proteins that were identified in <u>one of</u> <u>our recent publications</u> as DOP-interacting partners directly from brain tissues," Gendron said. "By studying their impact on the specific regulation of the trafficking and signaling of the DOP, we wish to eventually develop novel analgesics exhibiting fewer adverse effects than currently prescribed opioids."

More information: Jade Degrandmaison et al, Differential barcoding of opioid receptors trafficking, *Journal of Neuroscience Research* (2021). DOI: 10.1002/jnr.24949

Jade Degrandmaison et al, In vivo mapping of a GPCR interactome using knockin mice, *Proceedings of the National Academy of Sciences* (2020). DOI: 10.1073/pnas.1917906117

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