dioxygenase activity, GO:0051213

histone demethylase activity, GO:0032452 neural tube development, GO:0021915 cellular oxidant detoxification, GO:0098869 prostaglandin metabolic process, GO:0006693 histone H3-K9 demethylation, GO:0033169 leukotriene metabolic process, GO:0006691 prostaglandin biosynthetic process, GO:0001516 regulation of mRNA splicing, GO:0048024 histone demethylase activity (H3-K9 specific), GO:0032454 arachidonic acid metabolic process, GO:0019369 NAD biosynthesis via nicotinamide riboside salvage pathway, GO:0034356 respiratory system process, GO:0003016 hormone-mediated signaling pathway, GO:0009755 negative regulation of interleukin-10 production, GO:0032693 monocyte differentiation, GO:0030224 temperature homeostasis, GO:0001659 embryonic camera-type eye morphogenesis, GO:0048596 retinoic acid metabolic process, GO:0042573 retinal metabolic process, GO:0042574