

The orphan nuclear receptor Nurr1 is responsive to non-steroidal anti-inflammatory drugs

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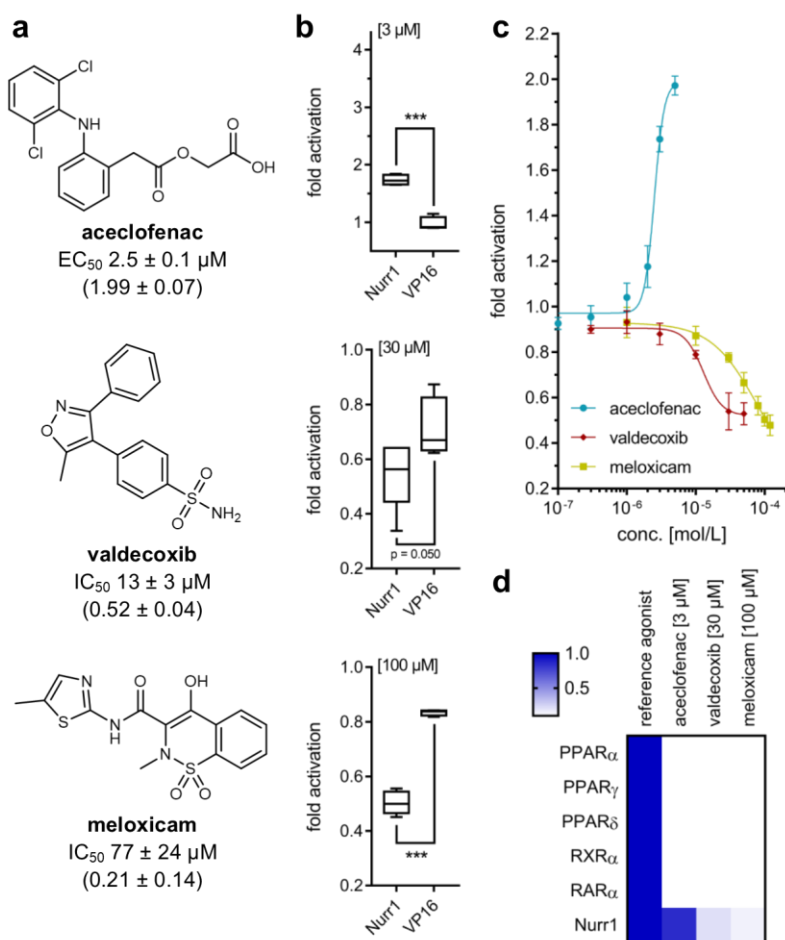
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- Supplementary Information -

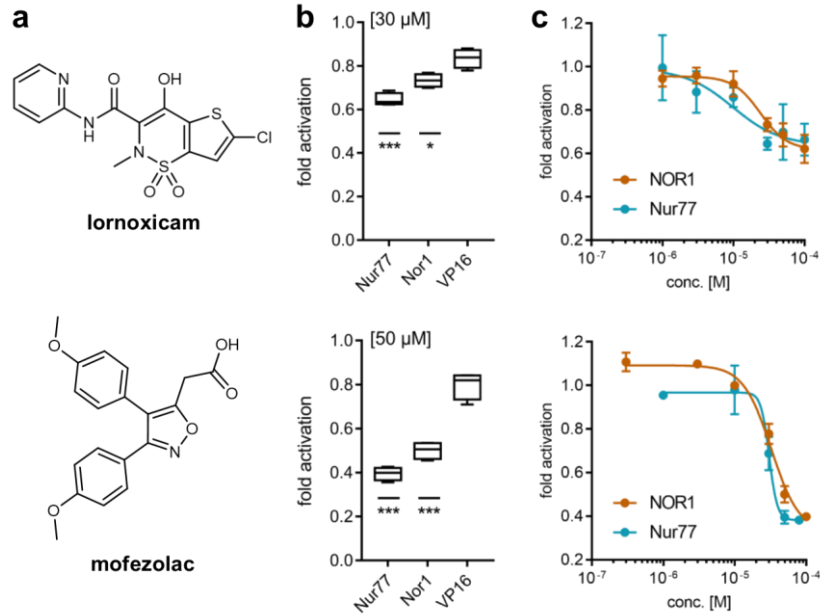
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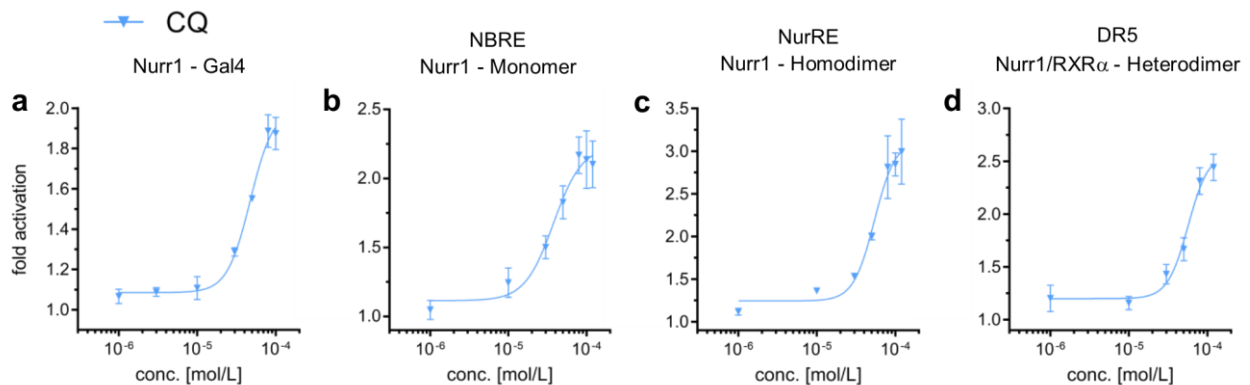
Supplementary Figures and Tables



Supplementary Figure 1. Bidirectional modulation of Nurr1 activity by drug approved COX inhibitors. (a) Molecular structures and activities of Nurr1 modulators aceclofenac, valdecoxib and meloxicam. EC_{50} and IC_{50} values were determined in the Gal4-Nurr1 hybrid reporter gene assay and are the mean \pm SD; $n \geq 3$. (b) Control experiments employing a Gal4-VP16 hybrid receptor confirmed Nurr1 mediated activity of aceclofenac, valdecoxib and oxaprozin. Boxplots show: center line, median; box limits, upper and lower quartiles; whiskers, min/max; $n \geq 4$. *** $p < 0.001$. (c) Gal4-hybrid reporter gene assay demonstrated Nurr1 activation by aceclofenac as well as inverse Nurr1 agonism for valdecoxib and meloxicam. Results are mean \pm S.E.M.; $n \geq 3$. (d) Selectivity profile of Nurr1 modulators over lipid activated transcription factors. Heatmap shows mean rel. activation which refers to reference agonists at 1 μM for PPARs (α : GW7647; γ : rosiglitazone; δ : L165,041), RXR α (bexaroten), RAR α (tretinoin) and 100 μM for Nurr1 (AQ); $n \geq 4$.



Supplementary Figure 2. Inverse agonists of related NR4A receptors Nur77 and NOR1 not affecting Nurr1 activity. (a) Molecular structures of NR4A receptor modulators lornoxicam and mofezolac. (b) Control experiments employing a Gal4-VP16 hybrid receptor confirmed Nur77 and NOR1 mediated activity of lornoxicam and mofezolac. Boxplots show: center line, median; box limits, upper and lower quartiles; whiskers, min/max; $n \geq 4$. * $p < 0.05$, ** $p < 0.01$ *** $p < 0.001$. (c) Dose-response curves demonstrate dose-dependent inverse Nur77 and NOR1 agonism. Data are mean \pm SD, $n \geq 3$.



Supplementary Figure 3. Cellular profiling of Nurr1 modulator chloroquine (CQ). Data shown here are identical with Fig. 3a-d for CQ in the manuscript but y-axis scaling has been adapted here to depict the CQ dose-response. The Nurr1 activation efficacy of CQ is markedly lower compared to AQ. (a) Gal4-hybrid reporter gene assay demonstrated Nurr1 activation by CQ. (b-d) Nurr1 full-length reporter gene assays with the human Nurr1 response elements NBRE (Nurr1 monomer, b), NurRE (Nurr1 homodimer, c), and DR5 (Nurr1:RXR heterodimer, d) confirmed agonism of CQ. All cellular experiments were performed in transiently transfected HEK293T cells. Results are the mean \pm S.E.M.; $n \geq 3$.

Supplementary Table 1. Activity of NSAIDs on NR4A nuclear receptors Nur77 (NR4A1), Nurr1 (NR4A2) and NOR1 (NR4A3) determined in uniform Gal4-hybrid reporter gene assays in transiently transfected HEK293T cells. Activity was verified using Gal4-VP16^{1,2} as control and only compounds with statistically significant ($p < 0.05$) activity on the respective NR4A receptor versus VP16 control are reported as active. EC₅₀/IC₅₀ values are reported in [μ M]. Values in parentheses are min./max. activation compared to 0.1% DMSO serving as vehicle. Data are the mean \pm SD, $n \geq 3$.

	Nur77	Nurr1	NOR1
meclofenamic acid	EC ₅₀ 3.9 \pm 0.7 (3.3 \pm 0.4)	EC ₅₀ 4.7 \pm 0.1 (3.52 \pm 0.05)	EC ₅₀ 7.9 \pm 0.8 (5.5 \pm 0.6)
meloxicam	IC ₅₀ 73 \pm 2 (0.23 \pm 0.02)	IC ₅₀ 77 \pm 24 (0.2 \pm 0.1)	IC ₅₀ 84 \pm 17 (0.1 \pm 0.1)
lornoxicam	IC ₅₀ 9.4 \pm 6.3 (0.63 \pm 0.09)	-	IC ₅₀ 24 \pm 3 (0.62 \pm 0.02)
aceclofenac	-	EC ₅₀ 2.5 \pm 0.1 (1.99 \pm 0.07)	-
mofezolac	IC ₅₀ 30 \pm 1 (0.38 \pm 0.02)	-	IC ₅₀ 33 \pm 5 (0.32 \pm 0.10)
oxaprozin	IC ₅₀ 16 \pm 5 (0.2 \pm 0.1)	IC ₅₀ 40 \pm 6 (0.26 \pm 0.08)	IC ₅₀ 22 \pm 4 (\geq 0.00)
valdecoxib	-	IC ₅₀ 13 \pm 3 (0.52 \pm 0.04)	-
parecoxib	IC ₅₀ 23.7 \pm 0.2 (0.1 \pm 0.0)	IC ₅₀ 13.4 \pm 0.3 (0.48 \pm 0.01)	IC ₅₀ 25 \pm 4 (0.22 \pm 0.05)

Supplementary References

1. Sadowski, I., Ma, J., Triezenberg, S. & Ptashne, M. GAL4-VP16 is an unusually potent transcriptional activator. *Nature* **335**, 563–564 (1988).
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