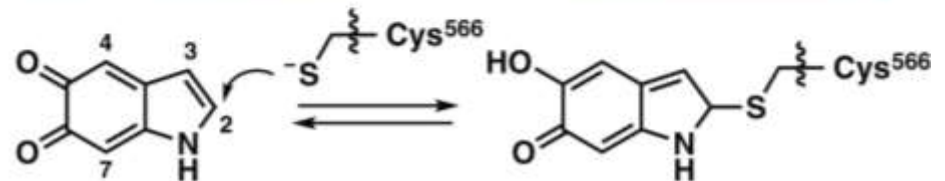
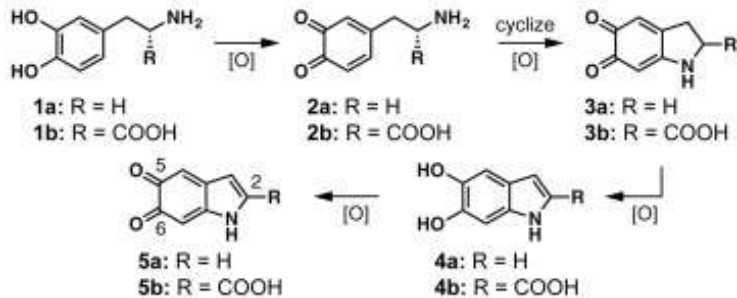
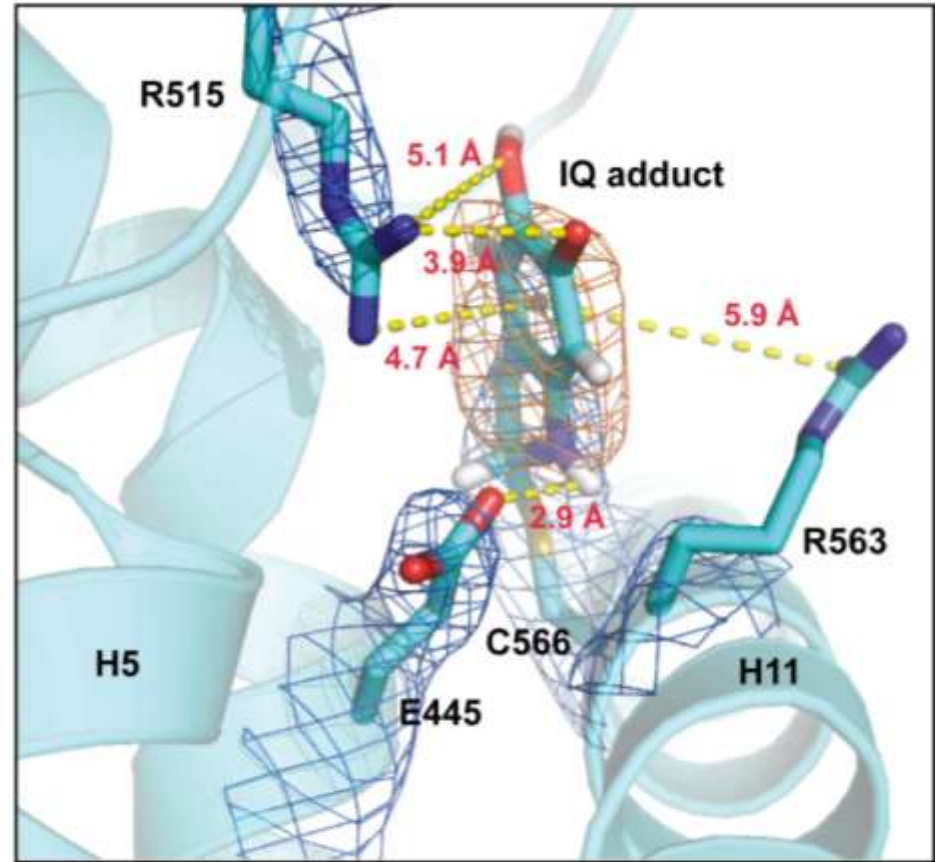
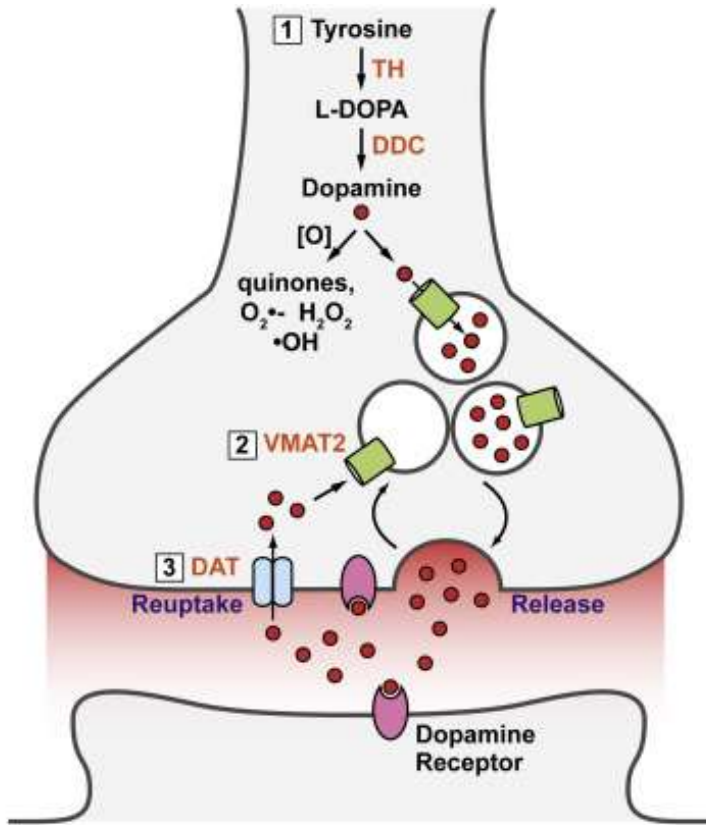


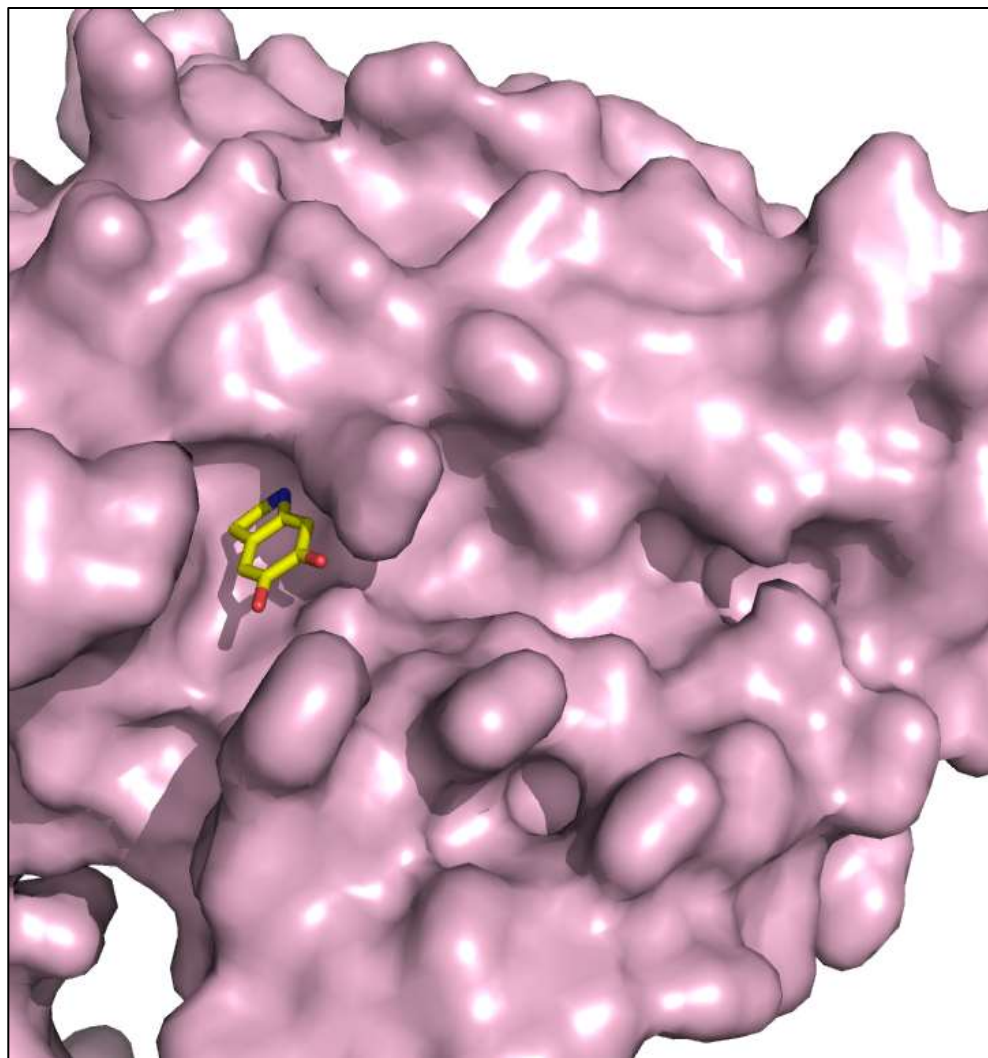
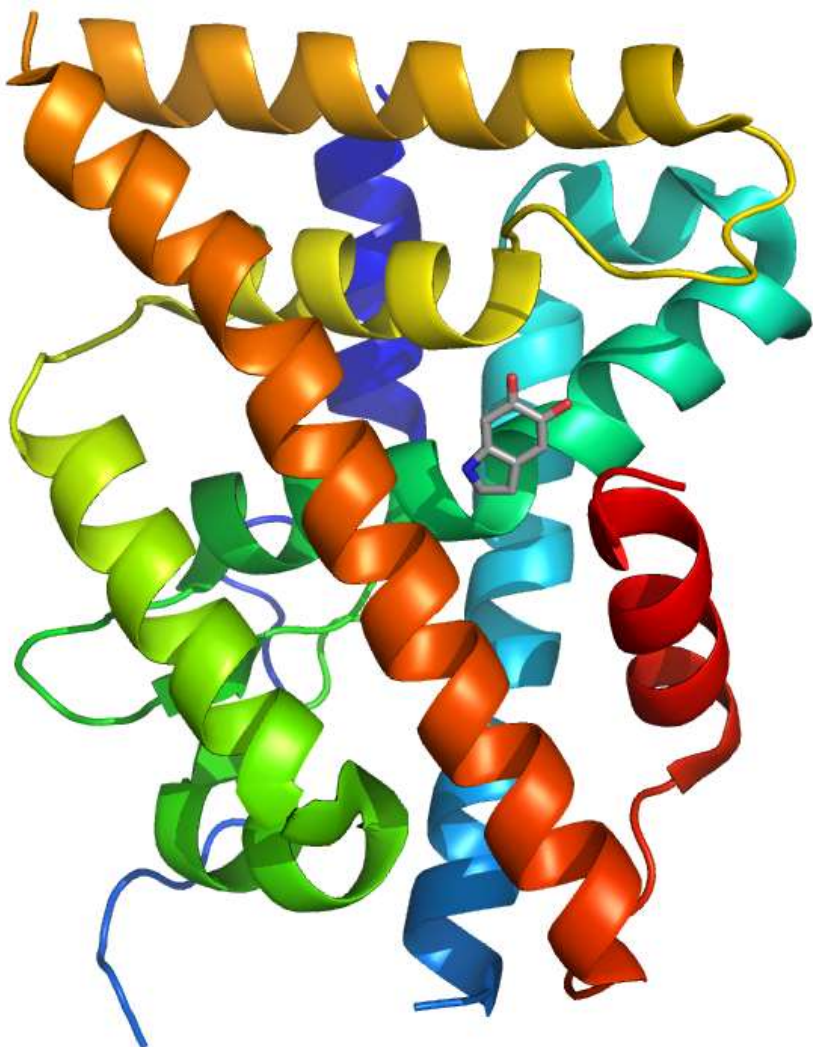
Structur-Guided Design of Nurr1 Agonists Derived from the Natural Ligand Dihydroxyindole

*J. Med.Chem.*2023,66,13556-13567

INTRODUCTION



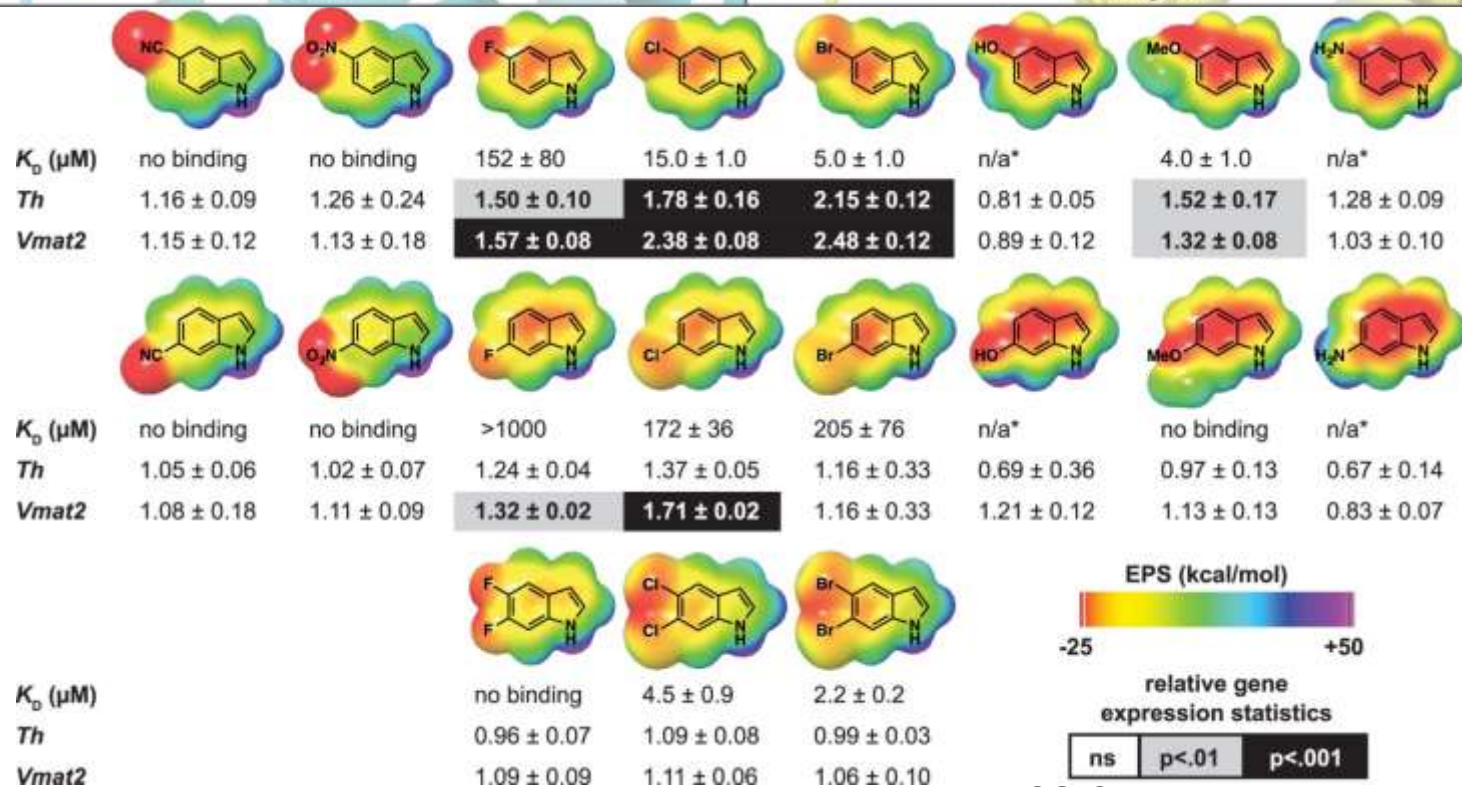
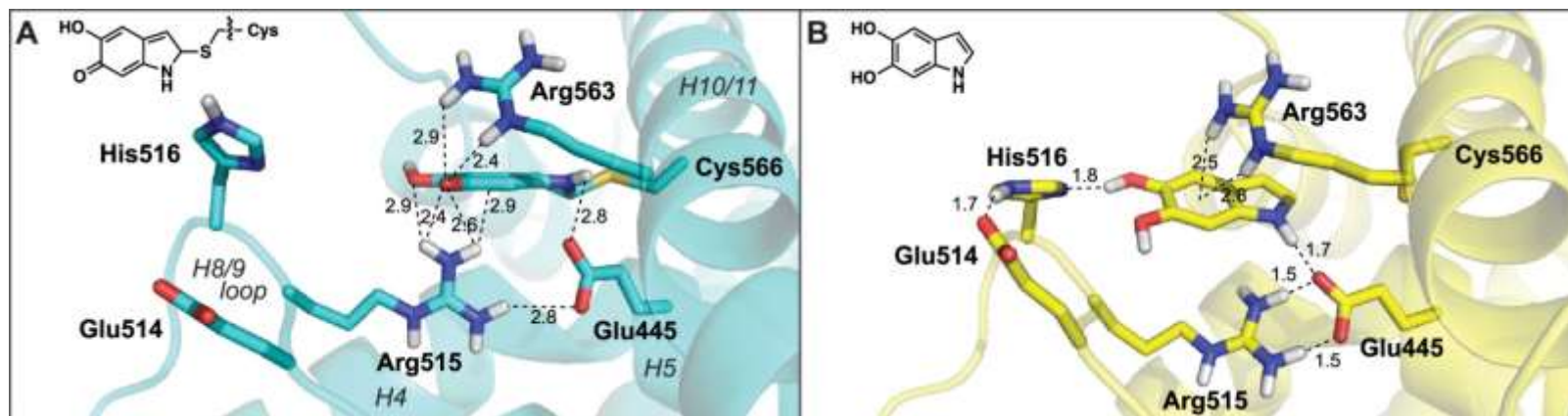
INTRODUCTION



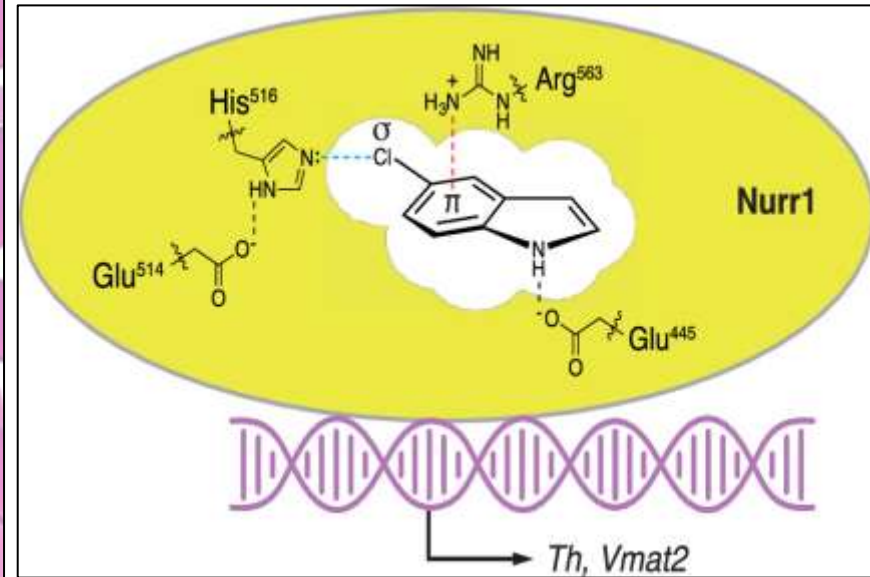
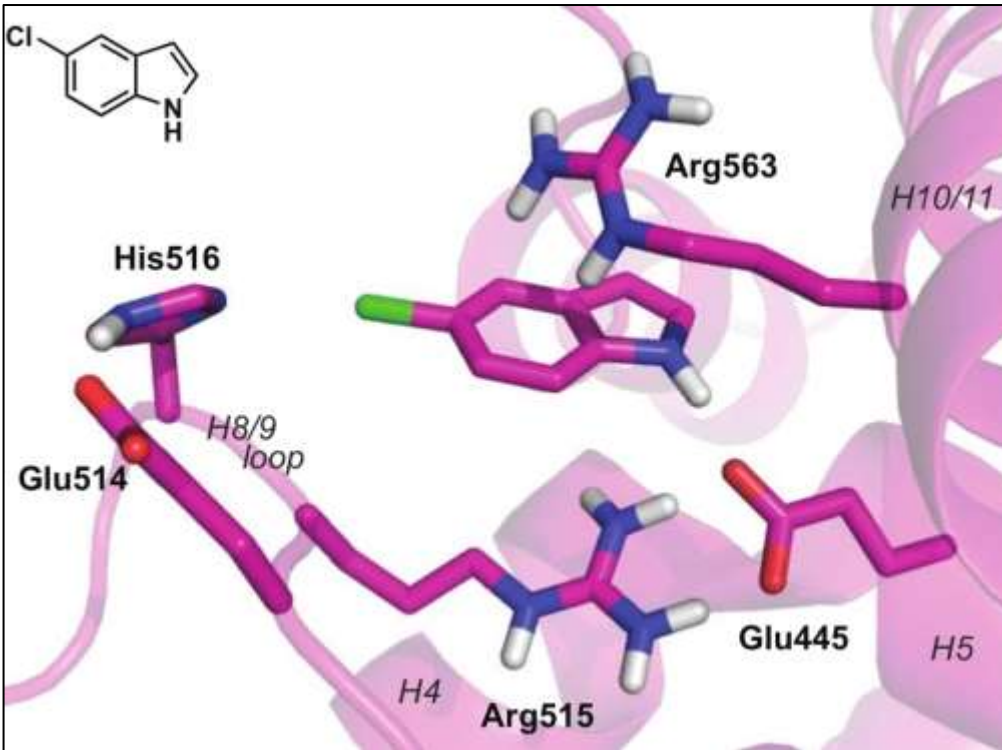
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Bruning JM. et al. *Cell Chem Biol.* 2019 May 16;26(5):674-685.e6.

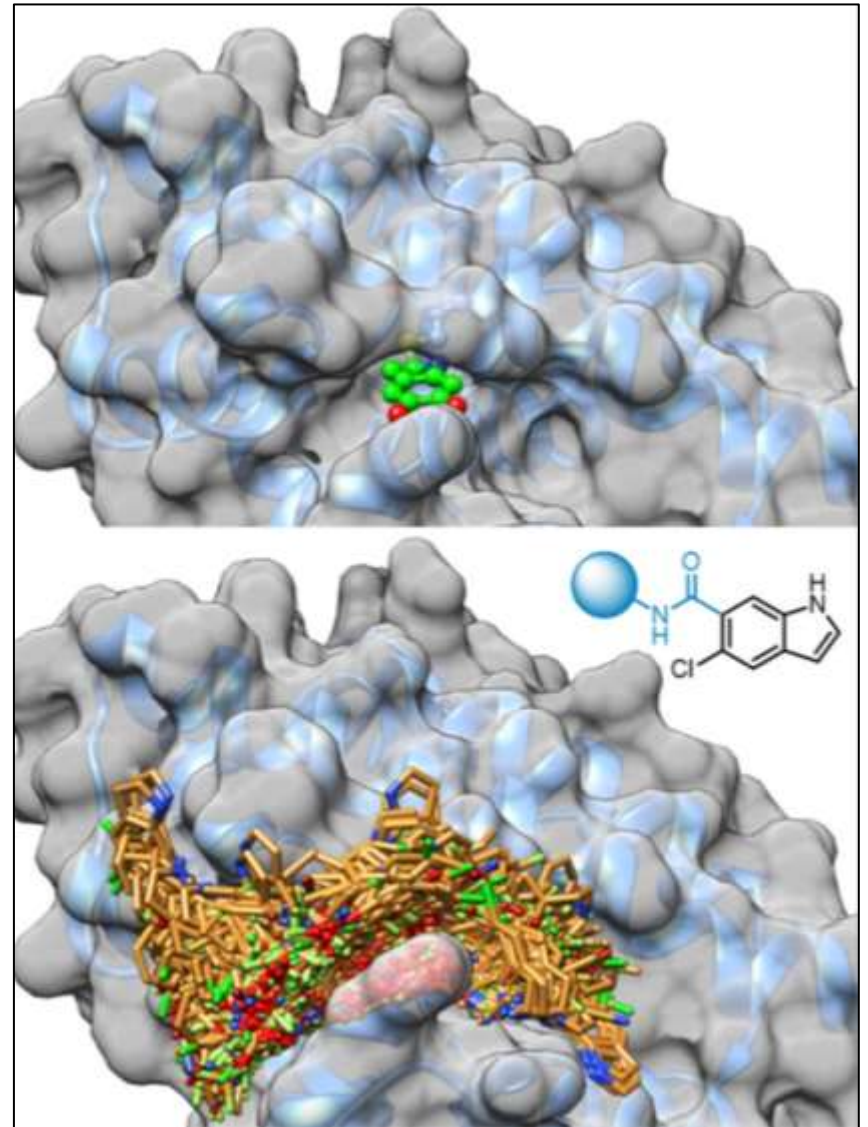
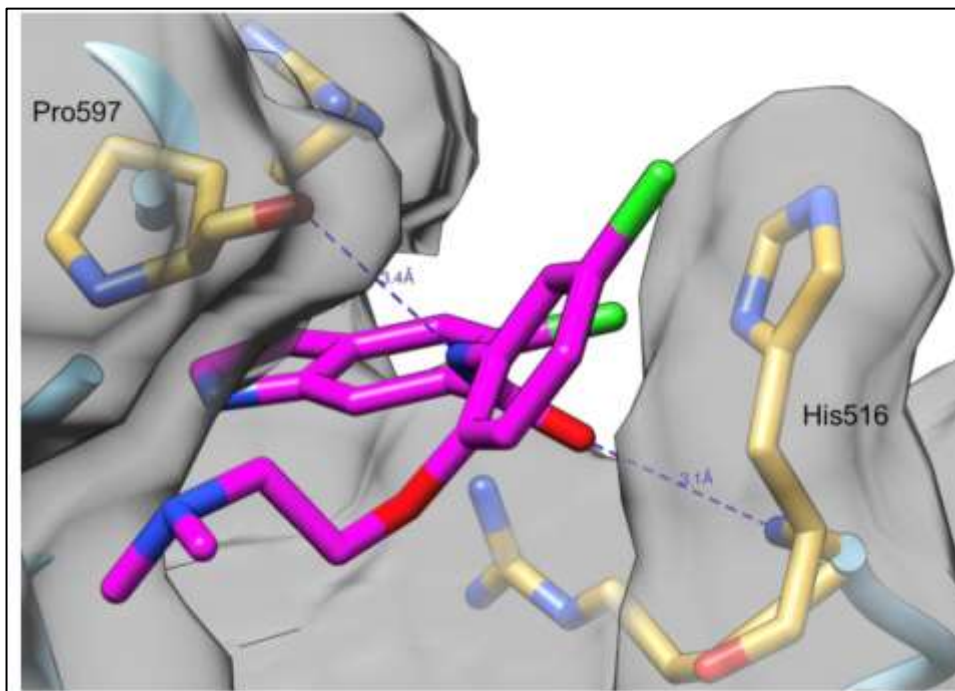
INTRODUCTION



Structure-Guided Design

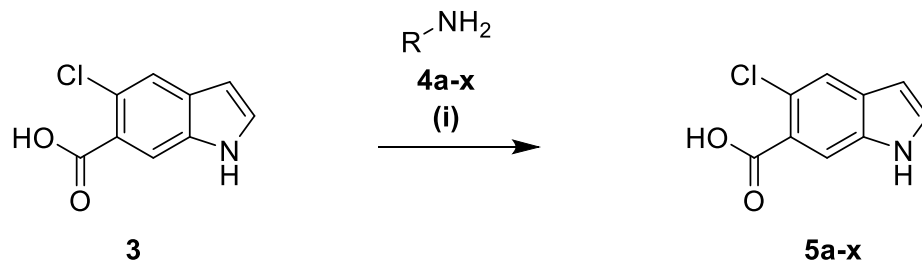


Structure-Guided Design

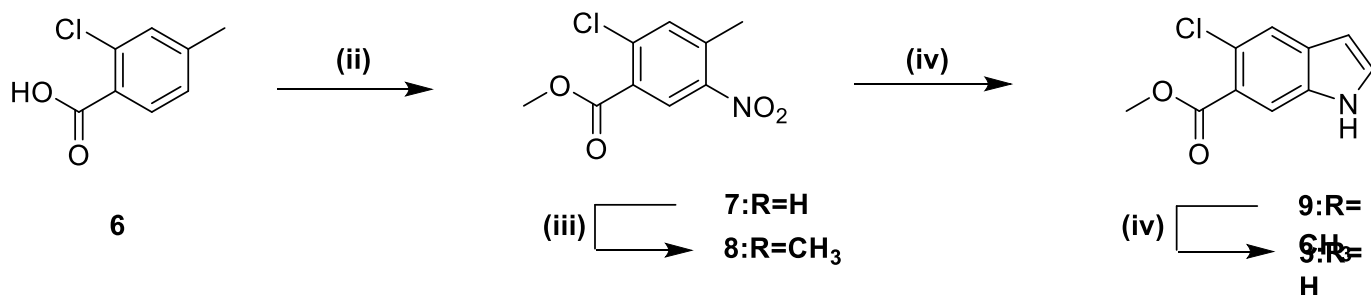


Compound Library Preparation and Screening

a



b



Reagents and conditions:

(i) EDC·HCl, EtOAc, rt, 36 h;

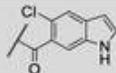
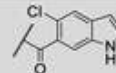
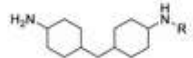
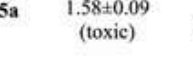
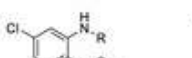
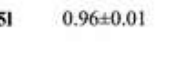
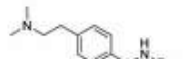
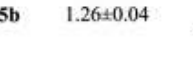
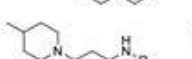
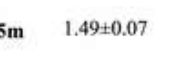
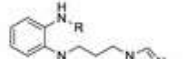
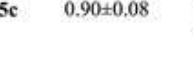
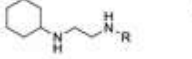
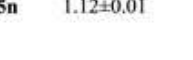
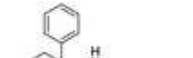
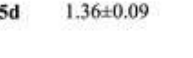
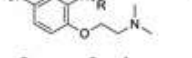

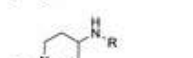
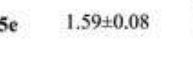


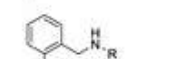
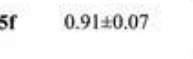
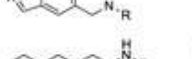
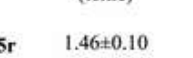

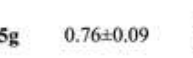
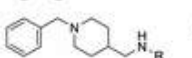
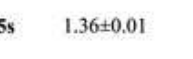

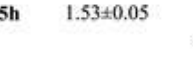
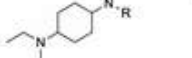


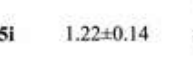
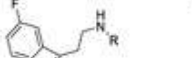
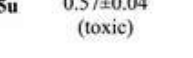
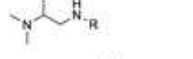

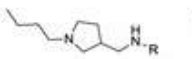
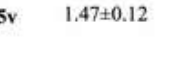
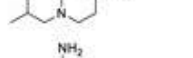
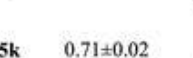
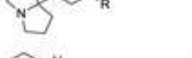


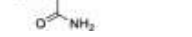

(ii) HNO₃/H₂SO₄, 5 °C, 0.5 h, 59%;

(iii) acetyl chloride, MeOH, 50 °C, 4 h, 96%;

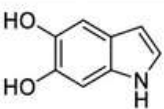
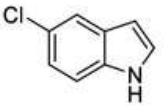
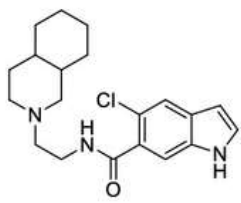
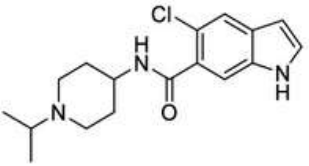
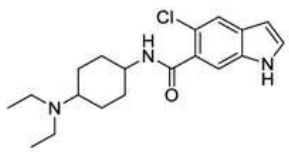
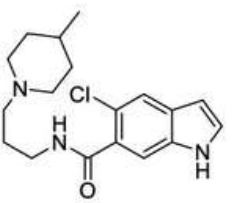
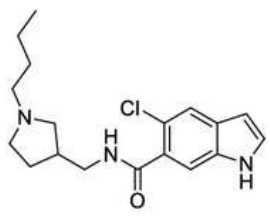
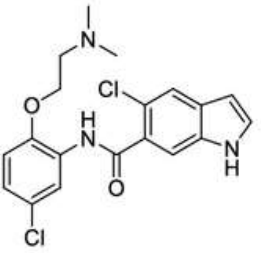
(iv) DMF-DMA, DMF, 120 °C, 2 h; then Zn, AcOH/H₂O, 80 °C, 2 h, 40%;

(v) LiOH·H₂O, EtOH/H₂O, rt, 18 h, 94%

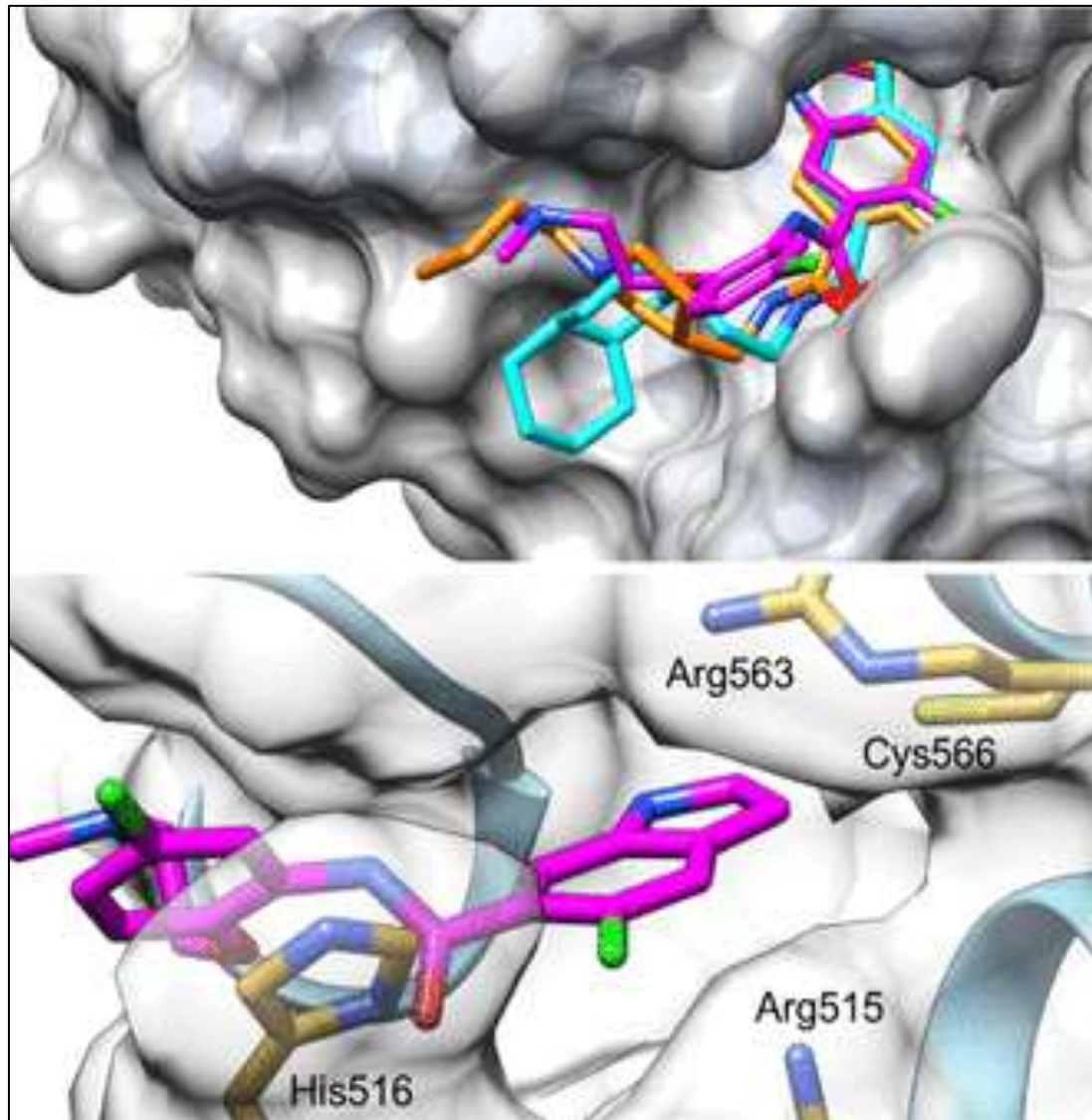
Compound Library Preparation and Screening

| R= -H | | R=  | | R= -H | | R=  | |
|---|--------------------|---|----------------------|--|----------------------|---|----------------------|
| ID | fold Nurr1 act. | ID | fold Nurr1 act. | ID | fold Nurr1 act. | ID | fold Nurr1 act. |
|  4a | 1.10±0.08 |  5a | 1.58±0.09 (toxic) |  4i | 0.89±0.07 |  5i | 0.96±0.01 |
|  4b | 1.62±0.15 |  5b | 1.26±0.04 |  4m | 1.08±0.04 |  5m | 1.49±0.07 |
|  4c | 1.29±0.09 |  5c | 0.90±0.08 |  4n | 0.96±0.12 |  5n | 1.12±0.01 |
|  4d | 1.60±0.14 |  5d | 1.36±0.09 |  4o | 1.19±0.05 |  5o | 1.60±0.01 |
|  4e | 1.10±0.12 |  5e | 1.59±0.08 |  4p | 1.51±0.19 (toxic) |  5p | 1.82±0.07 |
|  4f | 0.97±0.10 |  5f | 0.91±0.07 |  4q | 1.20±0.18 |  5q | 0.72±0.08 (toxic) |
|  4g | 1.24±0.05 |  5g | 0.76±0.09 |  4r | 1.05±0.13 |  5r | 1.46±0.10 |
|  4h | 1.6±0.4 (toxic) |  5h | 1.53±0.05 |  4s | 1.72±0.28 |  5s | 1.36±0.01 |
|  4i | 1.02±0.14 |  5i | 1.22±0.14 |  4t | 0.83±0.13 |  5t | 1.46±0.04 |
|  4j | 1.60±0.27 |  5j | 1.29±0.08 |  4u | 1.34±0.04 |  5u | 0.57±0.04 (toxic) |
|  4k | 1.09±0.05 |  5k | 0.71±0.02 (toxic) |  4v | 1.39±0.30 |  5v | 1.47±0.12 |
|  4l | | | |  4w | 0.70±0.15 |  5w | 1.26±0.04 |
| | | | | 4x | 0.86±0.08 | 5x | 0.89±0.06 |

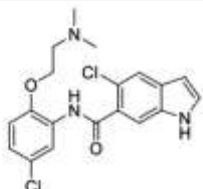
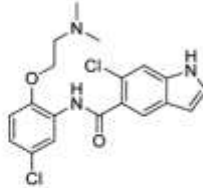
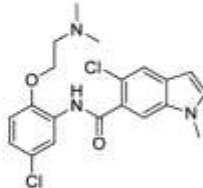
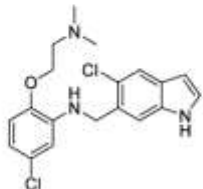
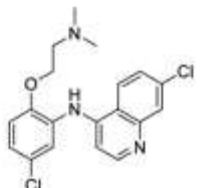
Compound Library Preparation and Screening

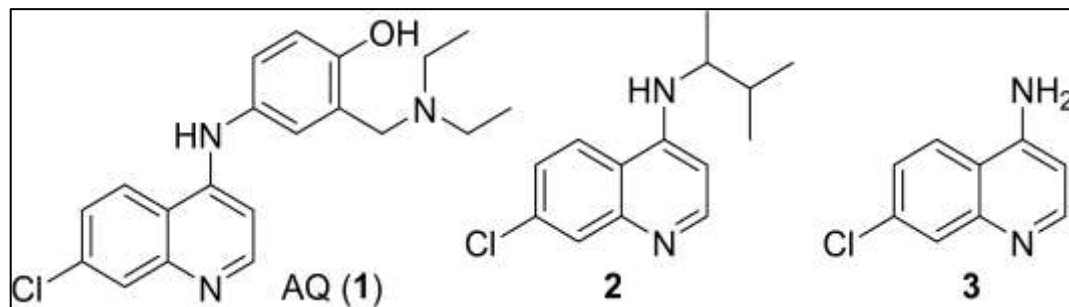
| ID | structure | K _d | EC ₅₀ (eff.) | | | |
|-------------|---|----------------------------|--------------------------------|----|--|---|
| DHI (2a) |  | - ^b | > 100 μM ²¹ | | | |
| 2b |  | 15 μM ²² | 40±4 μM (2.4±0.1-fold act.) | 5r |  | 3.2 μM 12±2 μM (1.4±0.1-fold act.) |
| 5e |  | no binding ^c | inactive at 100 μM | 5t |  | no binding ^c inactive at 100 μM |
| 5m |  | no binding ^c | inactive at 100 μM | 5v |  | 16 μM 28±6 μM (1.3±0.1-fold act.) |
| 5o |  | 0.5 μM | 3±1 μM (1.3±0.1-fold act.) | | | |

Binding Site Evaluation

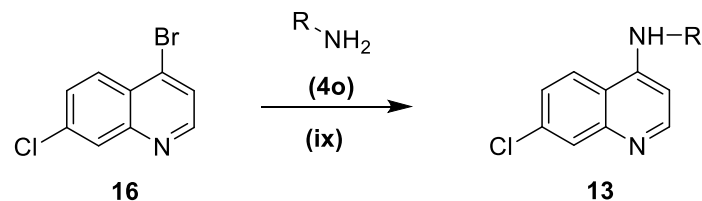
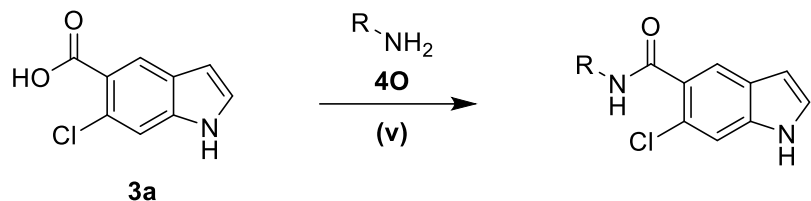
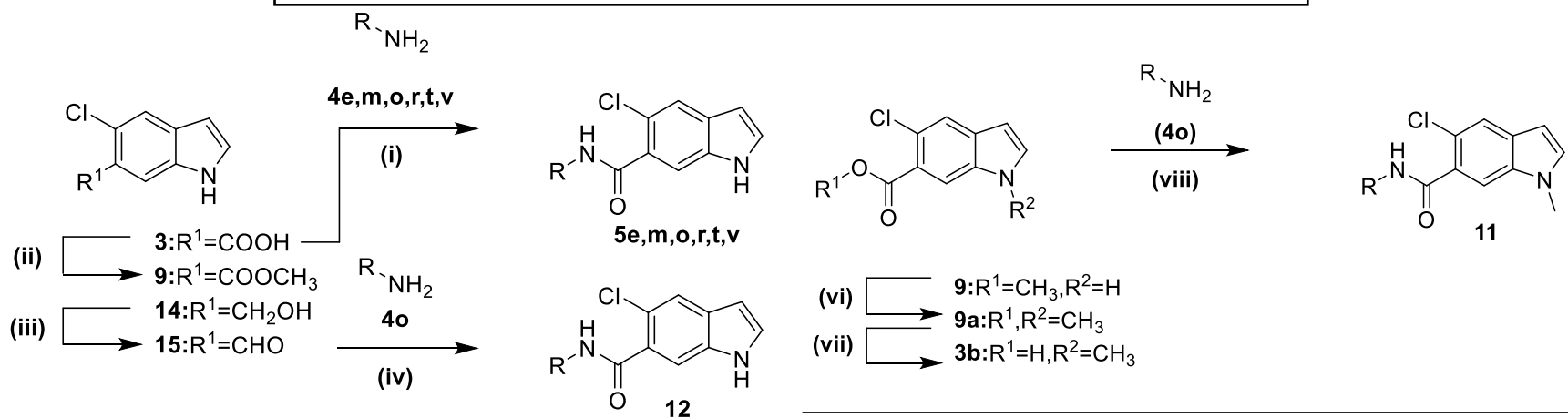


Structure-activity relationship

| ID | structure | K_d | EC_{50} (eff.) |
|----|---|-------------|---|
| 5o |  | 0.5 μ M | 3 \pm 1 μ M (1.3 \pm 0.1-fold act.) |
| 10 |  | 1.6 μ M | 6 \pm 3 μ M (1.4 \pm 0.1-fold act.) |
| 11 |  | 1 μ M | 16 \pm 6 μ M (1.4 \pm 0.1-fold act.) |
| 12 |  | 1.8 μ M | 5 \pm 2 μ M (1.4 \pm 0.1-fold act.) |
| 13 |  | 1.5 μ M | 3 \pm 1 μ M (1.5 \pm 0.1-fold act.) |

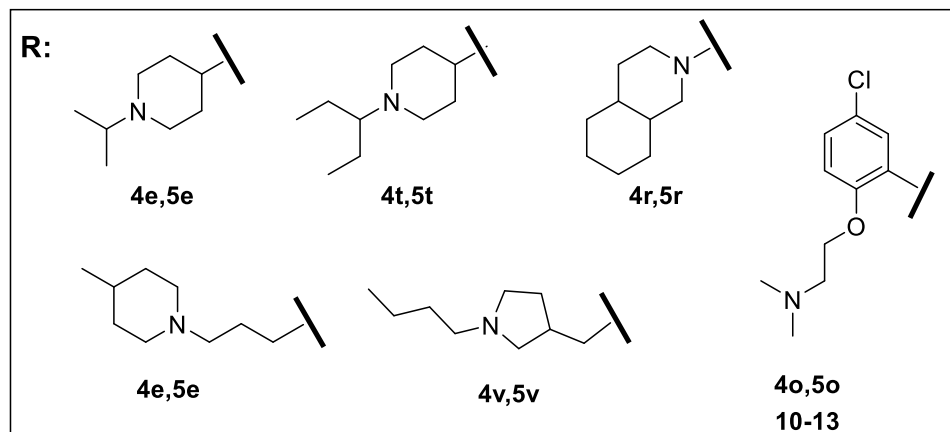


Structure-activity relationship

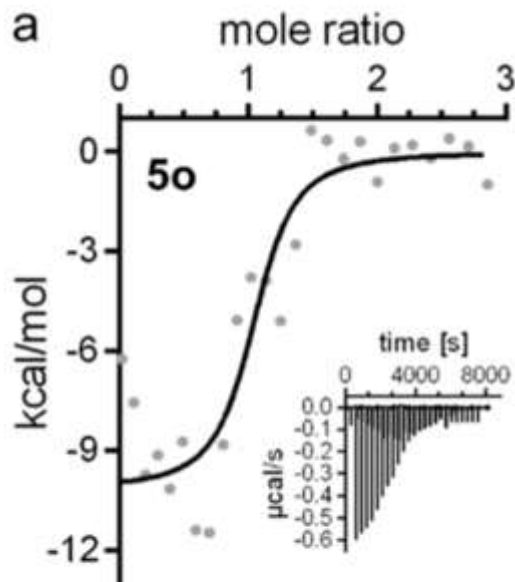


Reagents and conditions:

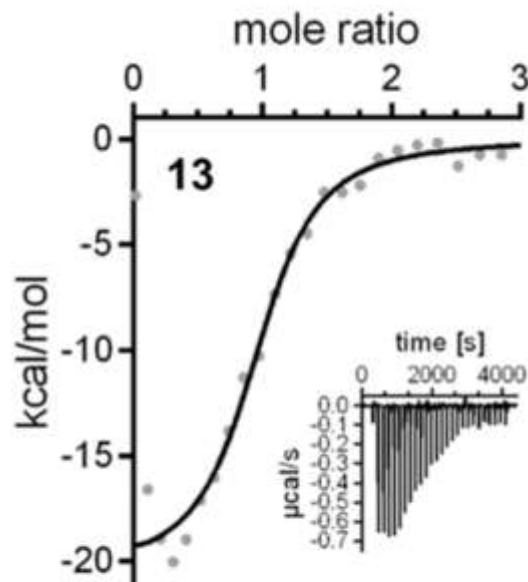
- (i) EDC·HCl, TEA, CHCl_3 , rt, 18 h, 8–32%;
- (ii) LiAlH_4 , THF, 0°C , 1 h, 71%;
- (iii) Dess–Martin periodinane, DCM, DMF, 0°C –rt, 1 h, 100%;
- (iv) $\text{NaBH}(\text{OAc})_3$, AcOH, DCM, DCE, rt, 2 h, 36%;
- (v) EDC·HCl, TEA, CHCl_3 , rt, 18 h, 5%;
- (vi) NaH, CH_3I , DMF, 0°C , 10 min, rt, 2 h, 33%;
- (vii) $\text{LiOH}\cdot\text{H}_2\text{O}$, EtOH, H_2O , rt, 18 h, 99%;
- (viii) NMI, TCFH, DMF, 80°C , 18 h, 26%;
- (ix) $\text{Pd}(\text{OAc})_2$, BINAP, K_3PO_4 , dioxane, 90°C , 24 h, 29%.



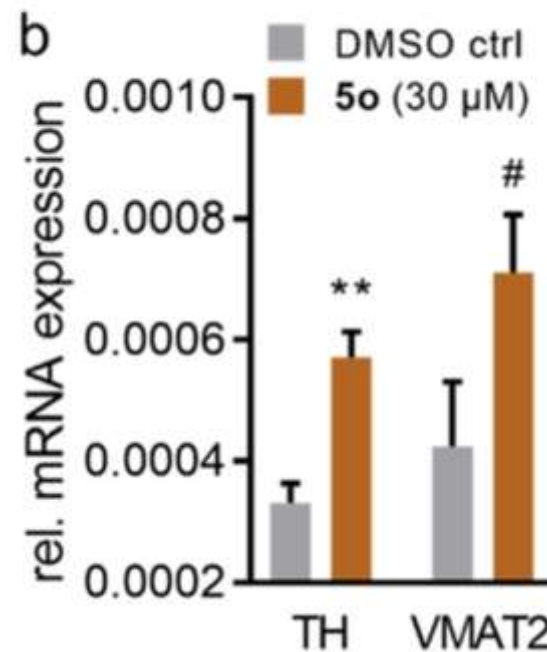
In vitro profiling of Nurr1 agonists



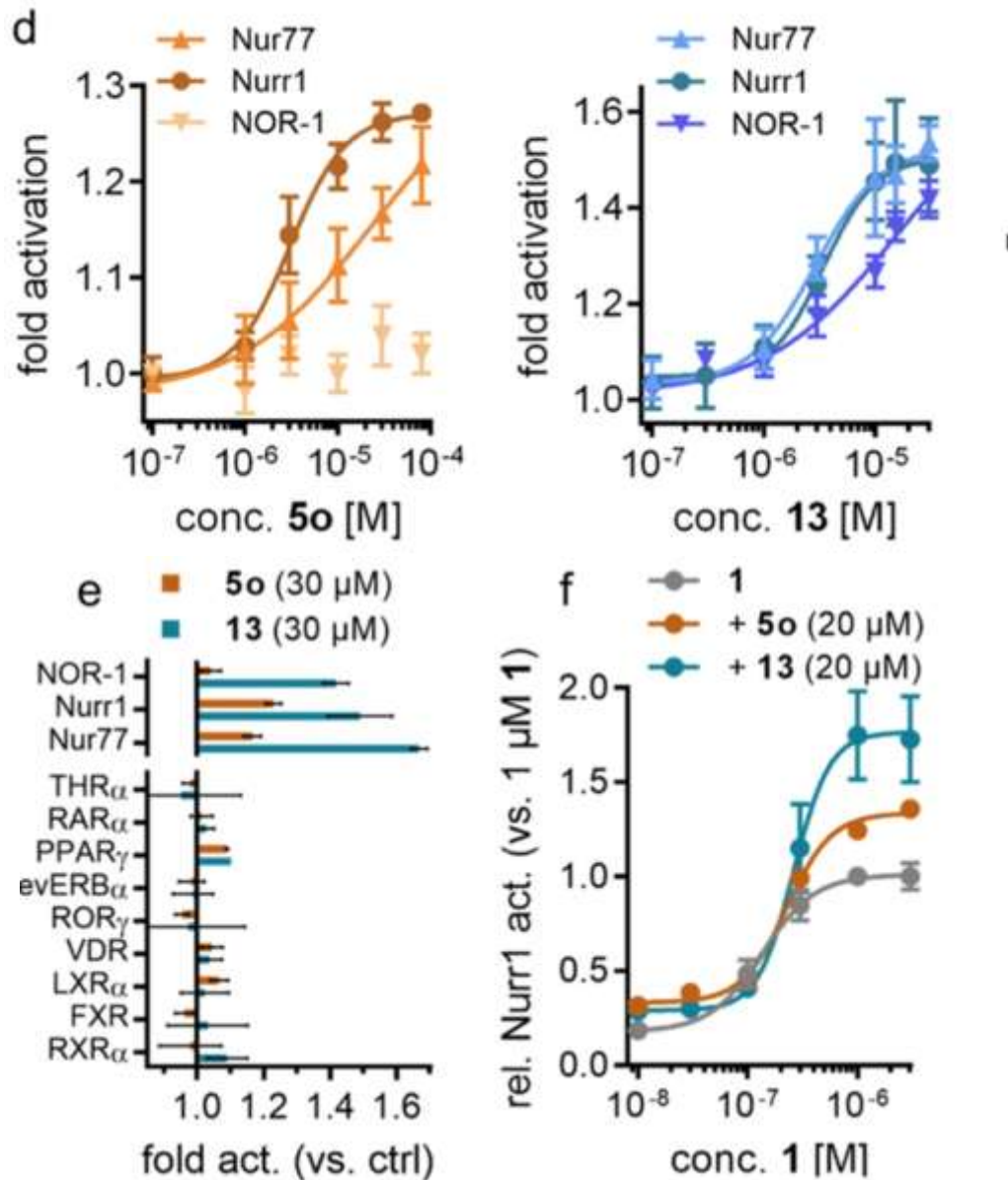
$K_d = 0.5 \mu\text{M}$, $n = 1.03$
 $\Delta H = -10.2 \text{ kcal/mol}$
 $-T\Delta S = 1.5 \text{ kcal/mol}$



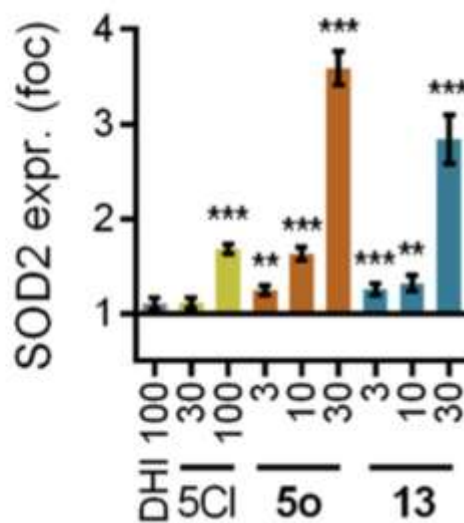
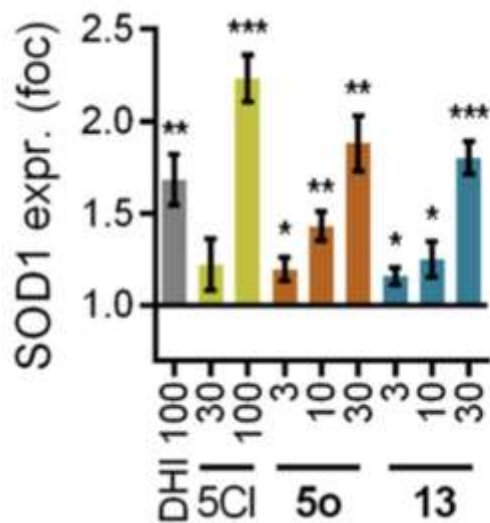
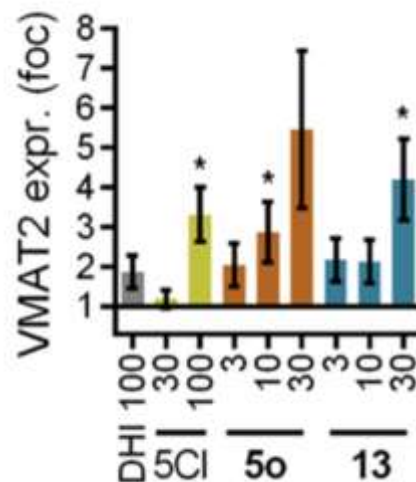
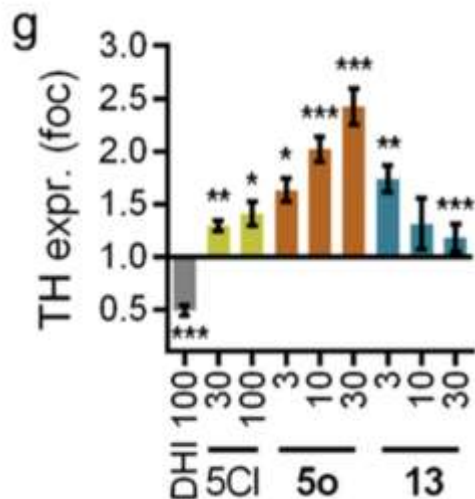
$K_d = 1.5 \mu\text{M}$, $n = 0.99$
 $\Delta H = -20.3 \text{ kcal/mol}$
 $-T\Delta S = 12.3 \text{ kcal/mol}$



In vitro profiling of Nurr1 agonists



In vitro profiling of Nurr1 agonists



CONCLUSIONS

- Drug discovery based on natural ligands of nuclear receptors has been very fruitful in the past as exemplified.
- The results demonstrate that the binding site of the natural (covalent) Nurr1 ligand DHI is druggable and can be addressed by non-covalent binders with markedly enhanced potency compared to the natural template, opening a new avenue to Nurr1 agonist development.
- 5o emerged from these DHI analogues with sub-micromolar affinity to Nurr1, preference over Nur77 and selectivity over NOR-1.
- Suggests that the binding site for Nurr1 is at the H12 helix, and despite some advances in the study of Nurr1 ligands, the understanding of this receptor is still very limited.

谢 谢 观 看

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