

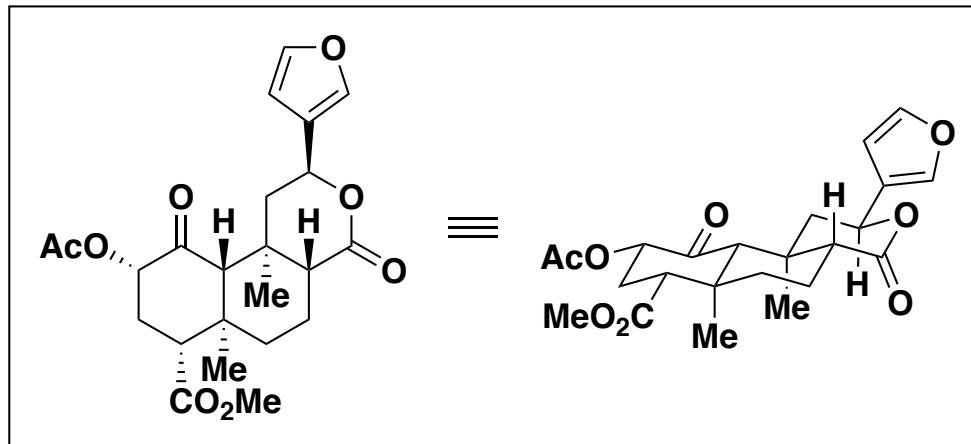
# ***Total synthesis of Salvinorin A***

Shenvi, R., Bohn, L. M. and coworkers.  
*ACS Cent. Sci.* **2017**, 3, 1329.

Wipf Research Group - Current Literature

Manwika Charaschanya  
August 11, 2018

# *Salvinorin A*

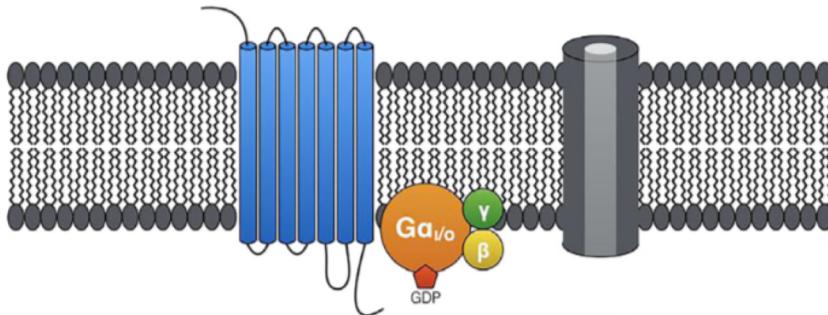


- Plant metabolite of *Salvia divinorum*
- Most potent naturally occurring hallucinogen
- Nonpeptide-like & non-nitrogenous scaffold (vs. opioid motifs)
- Diterpene with 7 chiral centers
- Agonist of *kappa*-opioid receptor (KOR), a subtype of G-protein-coupled receptor (GPCR)
- Selective for KOR over *mu*- and *delta*-opioid receptors (however, low affinity for MOR & shown to be an allosteric modulator of MOR)
- On-going race for biased KOR agonists towards clinical studies

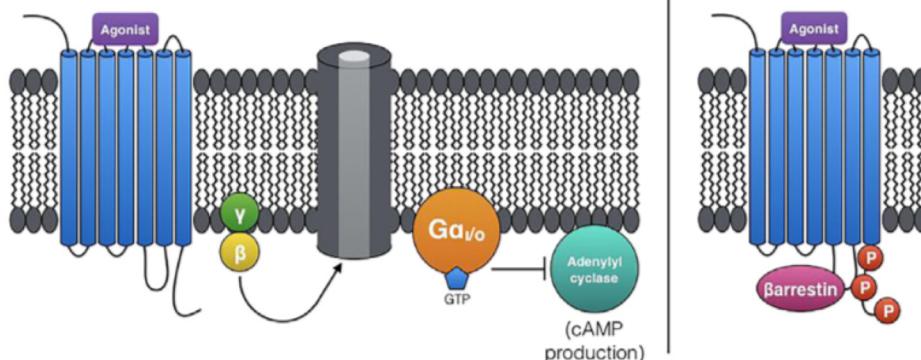
# Kappa opioid receptors

## GPCR signaling

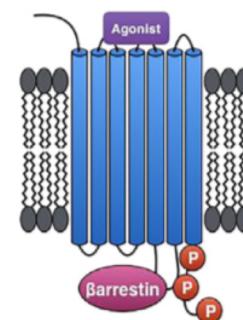
A. ORs in their unactivated state



B. G Protein Signaling



C. βarrestin Signaling



Balanced  
agonist



G protein  
signaling  
pathway  
βarrestin  
signaling  
pathway

analgesia  
hallucination  
sedation  
dysphoria  
diuresis

G protein  
biased  
agonist



G protein  
signaling  
pathway  
βarrestin  
signaling  
pathway

βarrestin  
biased  
agonist



G protein  
signaling  
pathway  
βarrestin  
signaling  
pathway

develop ligand  
for one  
physiological  
response

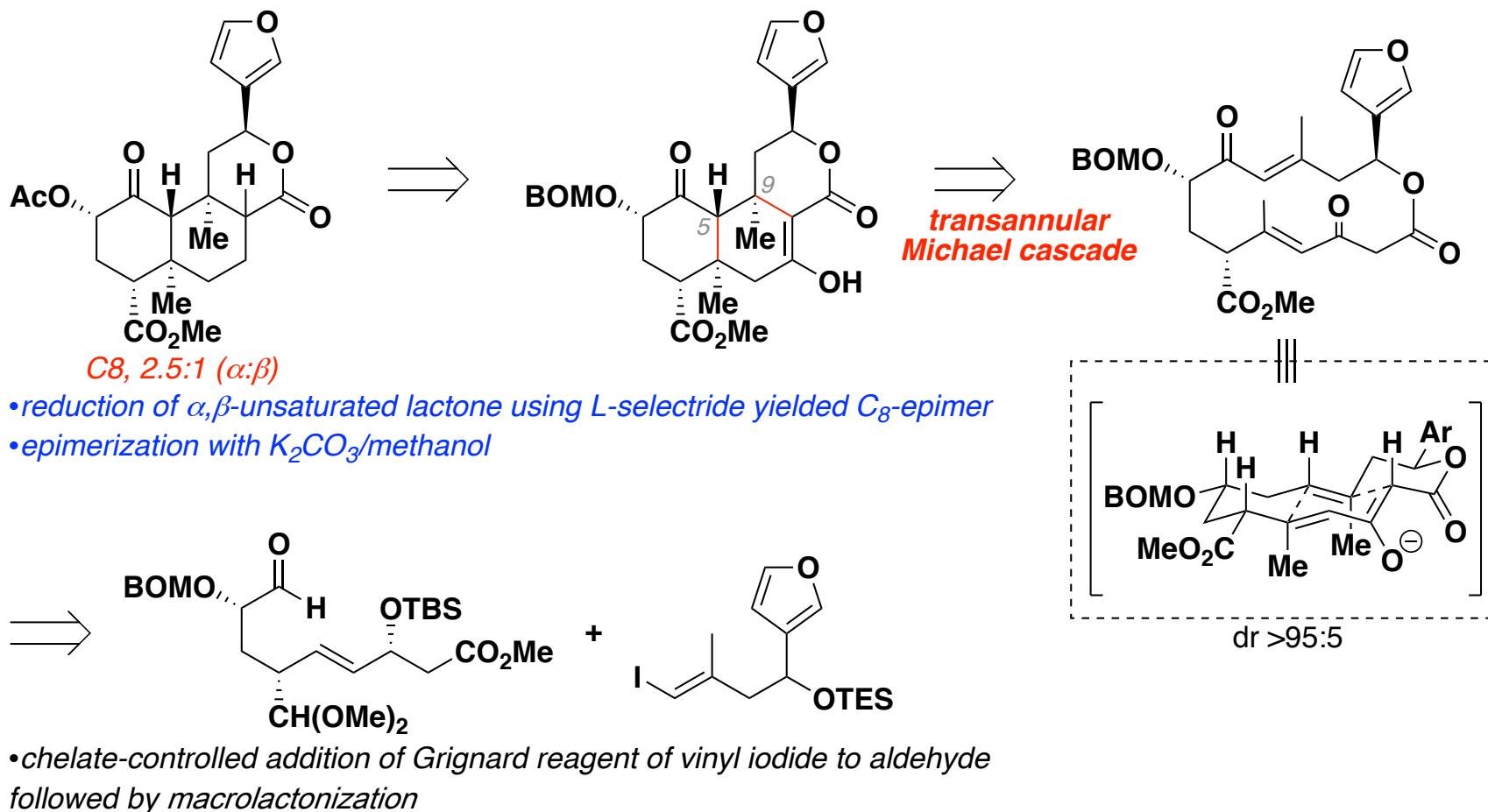
## ***Total syntheses of Sal A***

- First total synthesis by Rook (**2006**) was incomplete
- Evan's synthesis of Sal A was reported in 33 steps (**2007**)
- Hagiwara published a series of first- (**2008**) & second-generation syntheses (**2009**)
- Forsyth's synthesis of Sal A was reported in **2016**

# Total syntheses of Sal A

- Evan's synthesis of Sal A was reported in 33 steps (2007)

Retrosynthetic analysis:

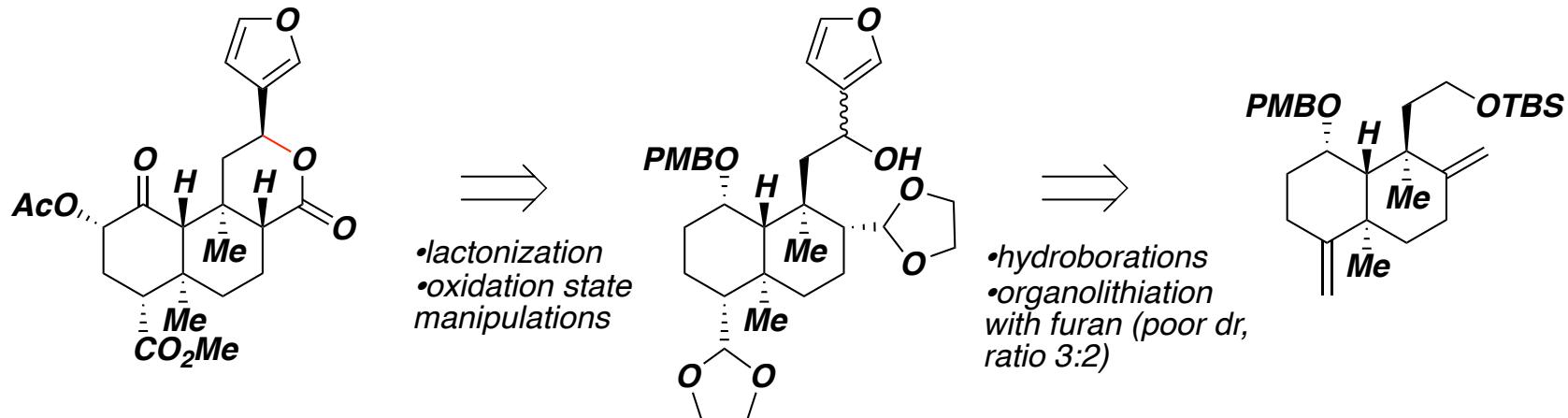


# Total syntheses of Sal A

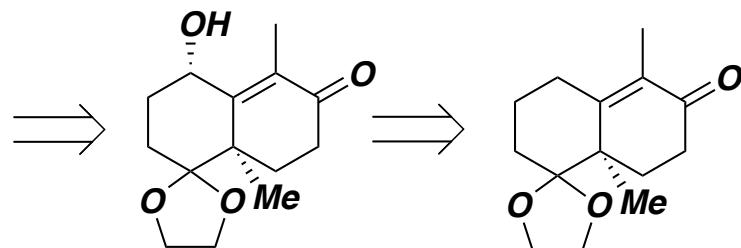
- Hagiwara first- (2008) & second-generation syntheses (2009)

## Retrosynthetic analysis:

### (a) First generation



•epimerization of C8 (dr 7:3); analogous to Evan's group



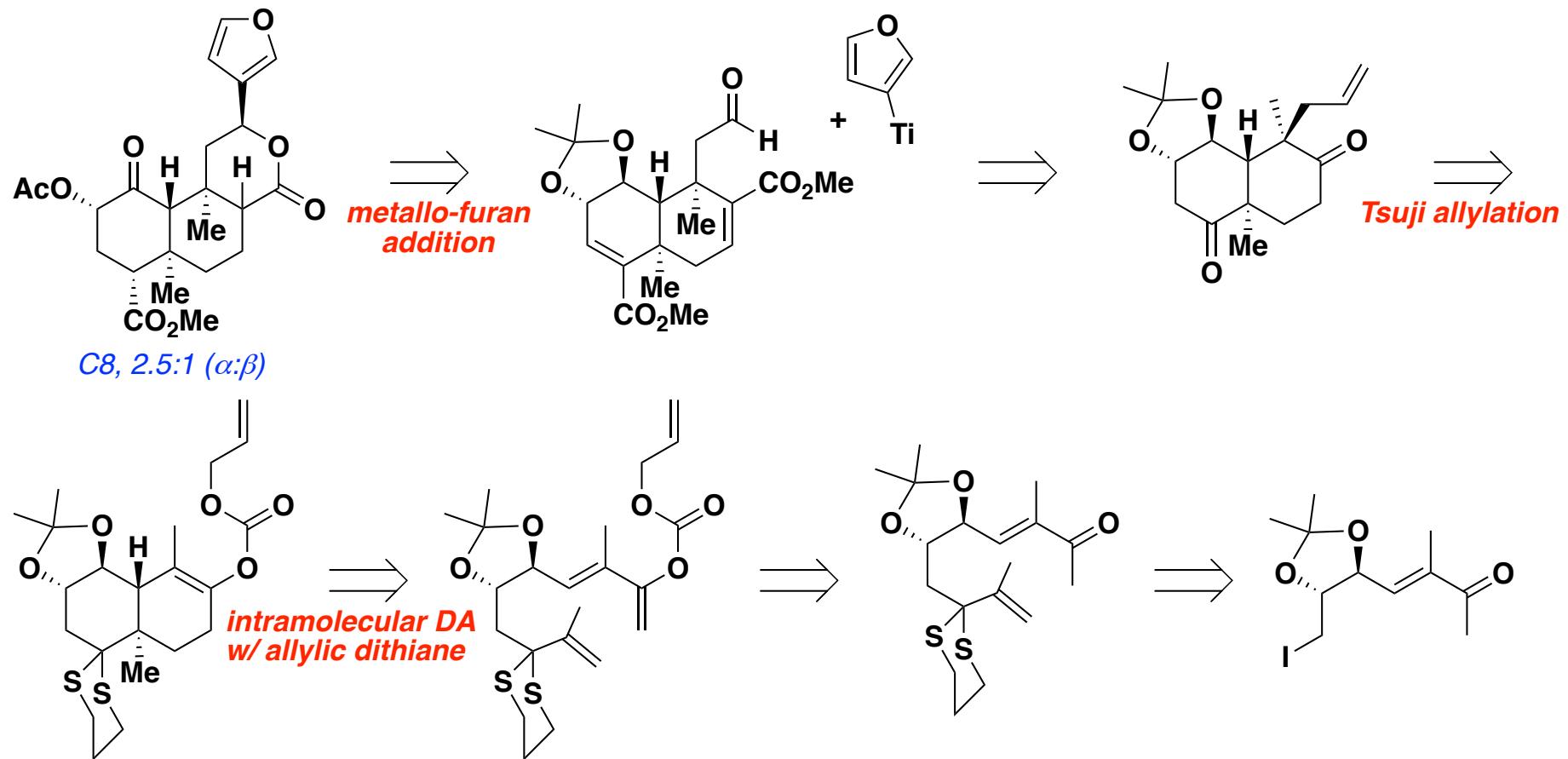
•originates from Wieland-Mischler ketone (known)

### (b) Second generation:

- resolved unfavorable selectivity of the 3-lithiofuran addition (use of furyl ketone vs. furyl alcohol)
- established shorter route; 20 steps (0.95% overall yield) to 13 steps (2.8% overall yield)

# Total syntheses of Sal A

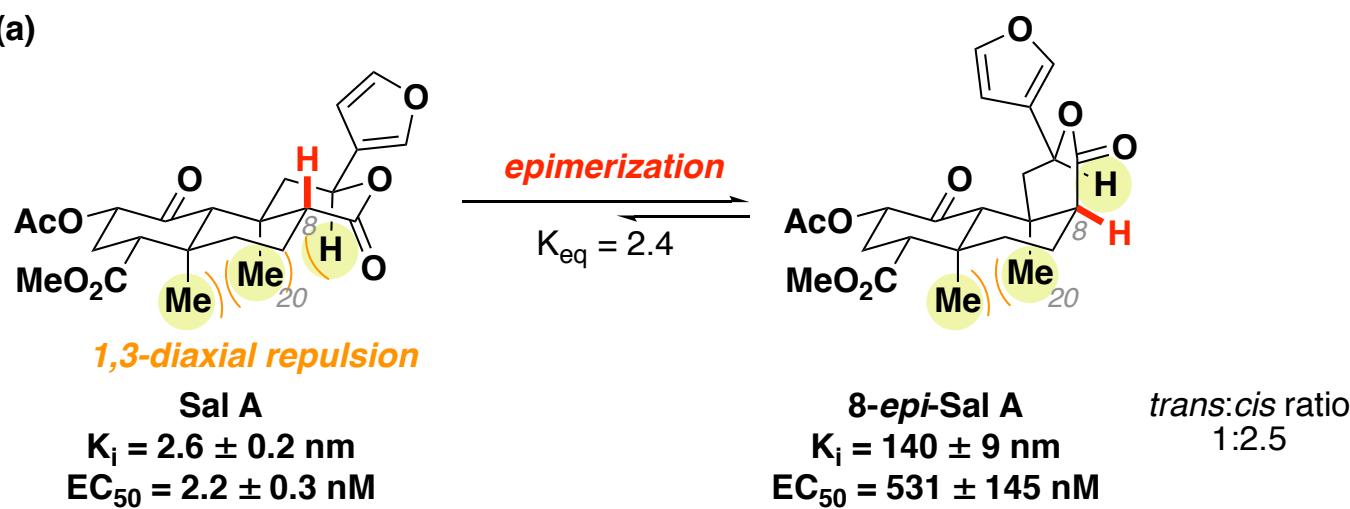
- Synthesis by Forsyth and coworkers (2016)



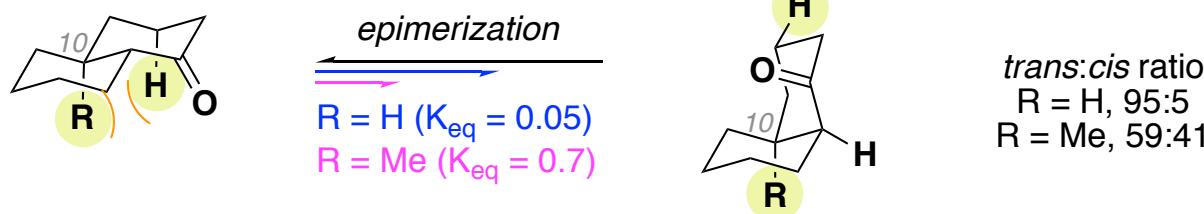
# Challenges in the total synthesis of Sal A

- Reactivity & thermodynamic stability of structural features: *trans*-decalin system, 6-membered lactone & furan
- Known configurational lability of C8 carbon – epimerization issue

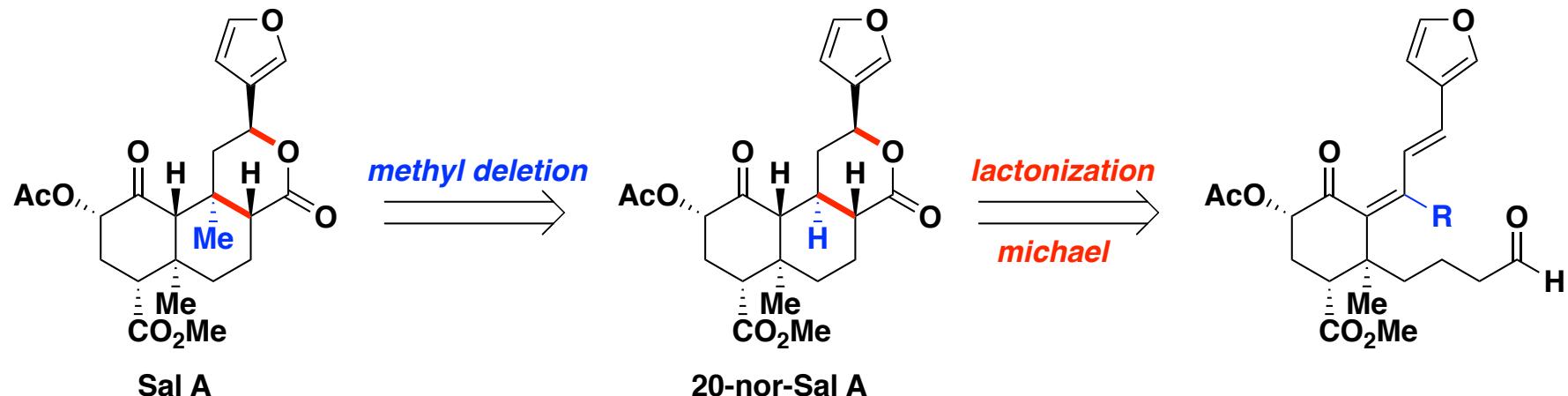
(a)



(b)



# *Dynamic strategic bond analysis yields 10-step synthesis of 20-nor-Sal A*

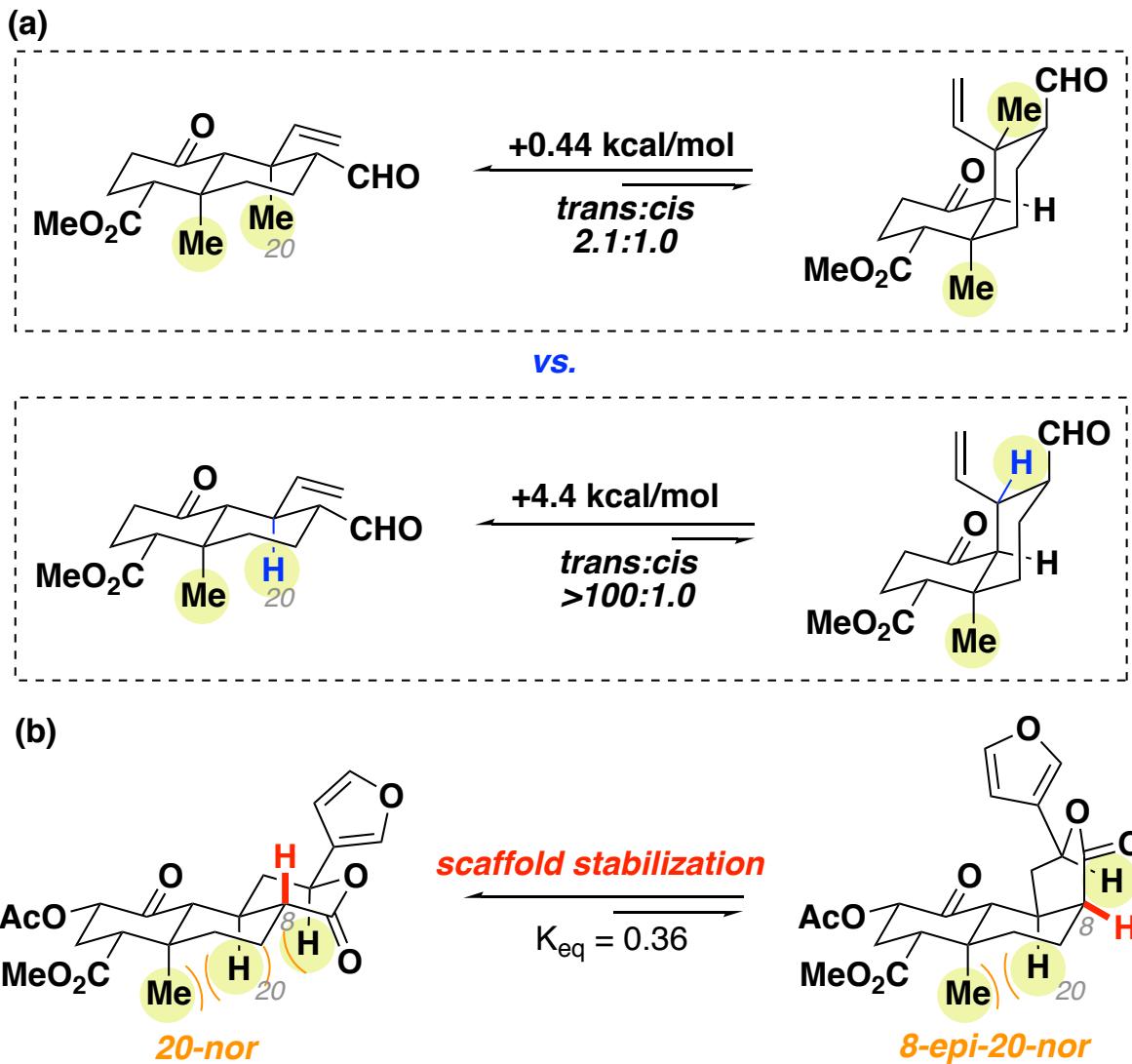


## Retrosynthetic analysis

- Hypothesis-driven
- Directly addresses known problem of configurational liability at C8
- Improve material access for med chem by 10-step synthesis

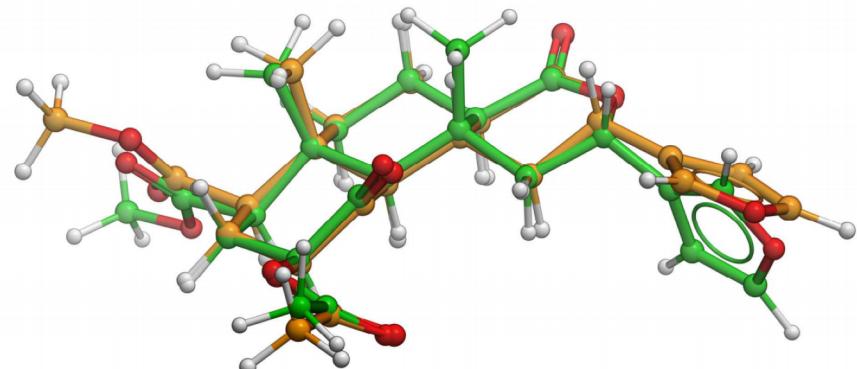
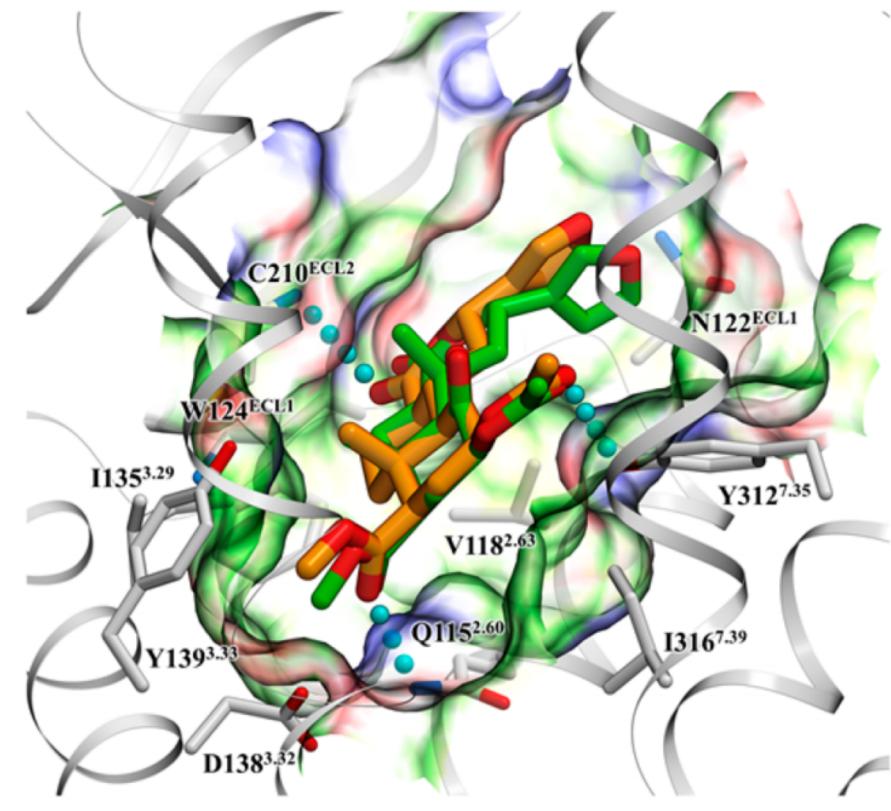
# Dynamic strategic bond analysis yields 10-step synthesis of 20-nor-Sal A

Hypothesis driven:

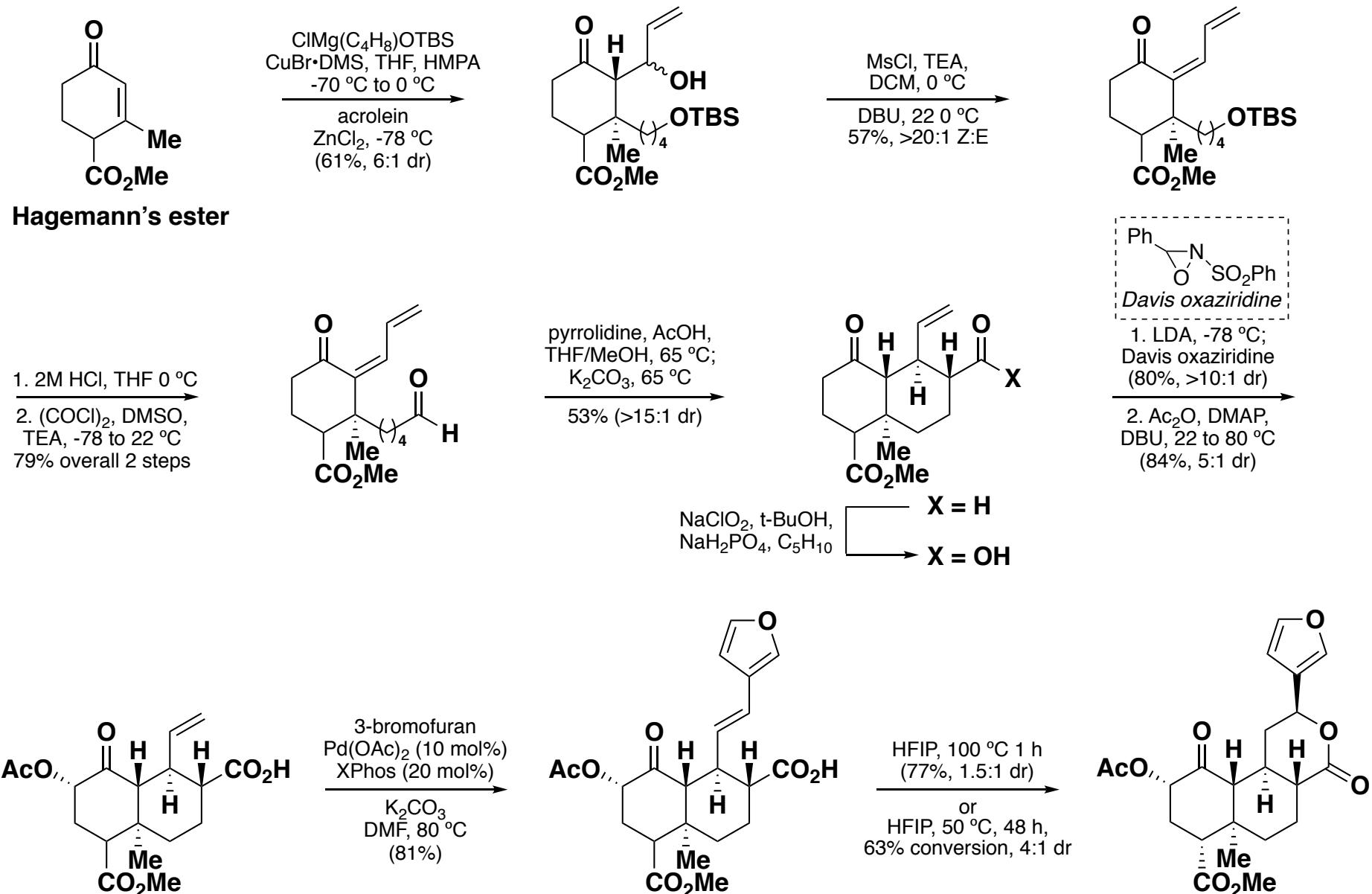


# *Docking mode of Sal A ligands*

- Sal A & 20-nor-Sal A bind in similar poses with comparable binding scores
- This binding pose suggested comparable binding affinity for Sal A & 20-nor-Sal A
- 20-methyl group directed toward extracellular region with no interactions with receptor
- Superimposition of structures show the geometry of the ring system maintained, however, removal of 20-methyl slightly displaces C-19 methyl and C-12 of lactone ring; resulting in subtle deviation in the position of the furan ring

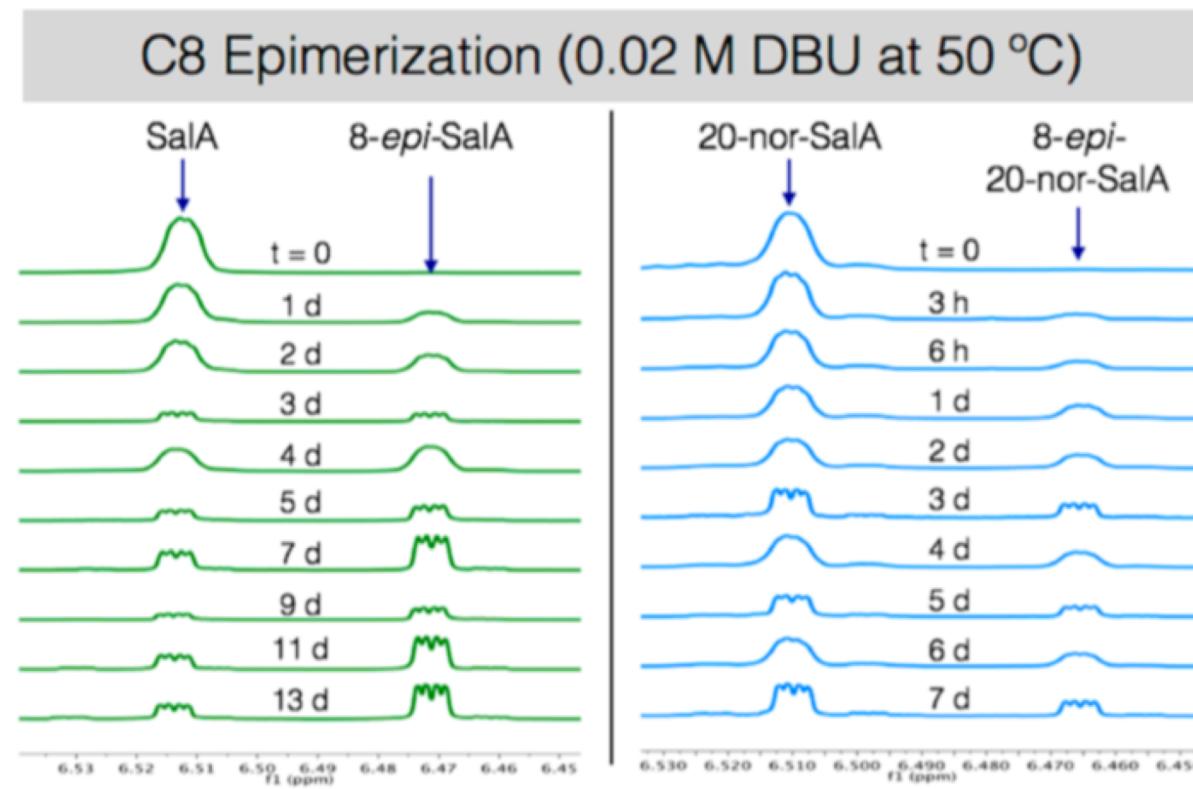


# Forward synthesis



# C8 Epimerization studies

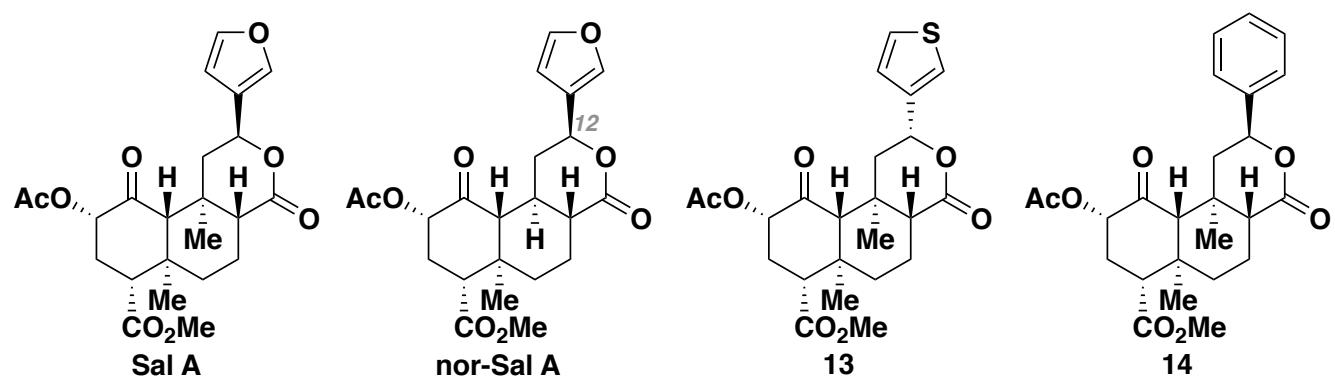
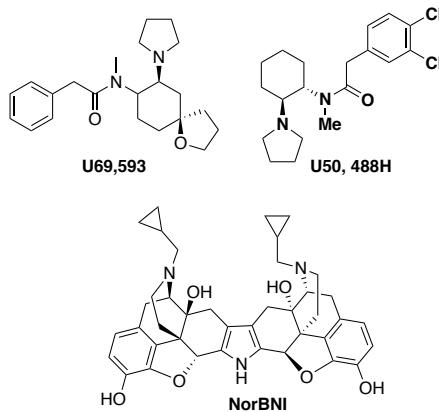
- 20-nor Sal A stabilizes the Sal A scaffold relative to its C8 epimer



- Follow-up paper (*Bioorg. Med. Chem.* 2018, *in press*): 06C-20-nor Sal A does not epimerize...even though ketone is more acidic than lactone

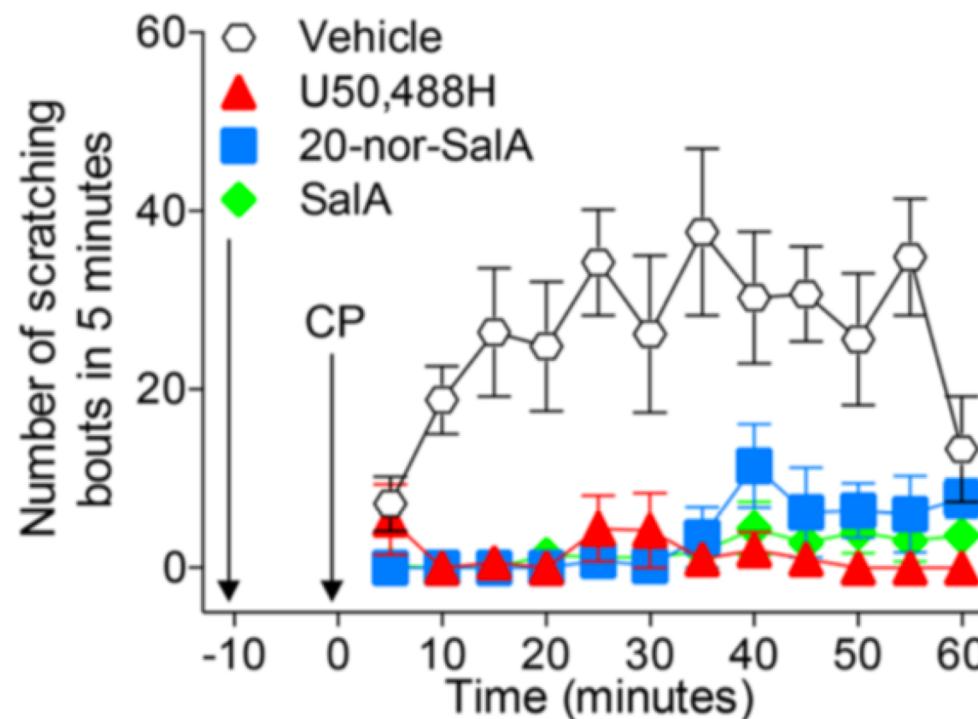
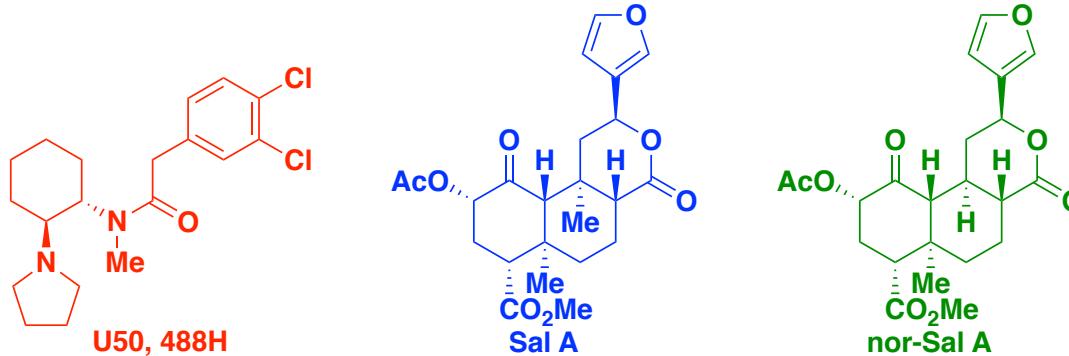
# Biological activity

Compound	Binding Affinities ( $K_i$ , nM)			$K_i$ ratios		Inhibition of cAMP	
	KOR	MOR	DOR	MOR/KOR	DOR/KOR	$EC_{50}$ , nM	% $E_{MAX}$
U69,593	1.07 ± 0.30	5176 ± 345	>10,000	4424	>153,177	5.7 ± 0.9	100
U50,488H	0.39 ± 0.24					7.5 ± 2.4	97 ± 1
NorBNI	0.04 ± 0.01					(antagonist)	
SalA	<b>0.16 ± 0.05</b>	616 ± 251	3417 ± 882	4741	26,285	<b>0.9 ± 0.3</b>	99 ± 1
20-nor-SalA	<b>1.08 ± 0.36</b>	7994 ± 2247	>10,000	7402	>10,933	<b>6.1 ± 1.7</b>	99 ± 2
12- <i>epi</i> -20-nor-1	<b>2.13 ± 1.20</b>	5113 ± 1547	3628 ± 913	2400	1703	<b>84 ± 13</b>	102 ± 1
13	<b>0.76 ± 0.25</b>	2841 ± 422	2367 ± 748	3738	3114	<b>23 ± 5.9</b>	100 ± 0
14	<b>0.53 ± 0.20</b>	4546 ± 516	>10,000	8577	>23,206	<b>16 ± 5.8</b>	95 ± 3
12- <i>epi</i> -14	<b>0.79 ± 0.22</b>	5683 ± 1543	5171 ± 633	7194	6546	<b>15 ± 3.5</b>	101 ± 1



# *Biological activity*

- Kappa agonists suppress chloroquine phosphate-induced pruritus in mice



# Summary

- Integration of *in silico* docking and retrosynthetic analysis can prompt scaffold redesign --- *dynamic retrosynthetic analysis* ---
  - ✓ maintained scaffold complexity w/ synthesis simplification
  - ✓ maintained target engagement w/ increased scaffold stability
- New opportunities to identify opioid receptor probes & drugs

