



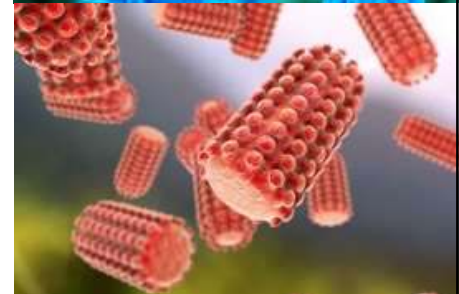
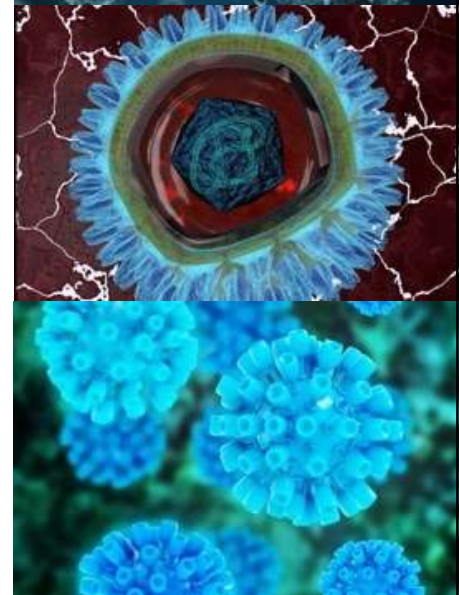
The Allosteric Sirtuin 2 Modulator FLS-359 Inhibits Viral Replication

Stacy Remiszewski, Ph.D.

Invited Speaker

Denver ACS Meeting August 19, 2024

Session: Approaches to the Modulation of Enzyme
Activation and/or Substrate Recognition



Agenda

- Introduction & identification of sirtuins (SIRT2s) as viral restriction factors
- SIRT2 Enzyme mechanism and overview of SIRT2 structure
- FLS-359 broad spectrum antiviral activity and SIRT2-mediated cellular effects
- Mechanism of FLS-359 SIRT2 enzyme activity modulation
- Comparison structures of FLS-359 & literature SIRT2 modulators
- Summary

Human cytomegalovirus (HCMV)

- δ b-herpesvirus: 200 ORFs and 28 miRNAs
- δ Ubiquitous: 1% of newborns; 10% 6-month-olds; 60-99% of adults
- δ Infection in healthy individuals generally asymptomatic
 - δ Acute phase (weeks) - many cell types & organs
 - δ Persistent phase (years) - salivary, mammary & kidney epithelial cells
 - δ Latent phase (lifetime) - CD34⁺ bone marrow cells
- δ Congenital infection: hearing loss, profound CNS damage
- δ Infection can present in immunosuppressed: AIDS & transplant
 - δ Provides an opportunity to develop a therapy for an unmet medical need
- δ Standards of care - direct acting antivirals: ganciclovir, letermovir, maribavir
 - δ DNA polymerase (pUL54) inhibitor, terminase (pUL56) inhibitor and kinase (pUL97) inhibitor

Direct acting antivirals (DAAs) suffer from therapeutic shortcomings

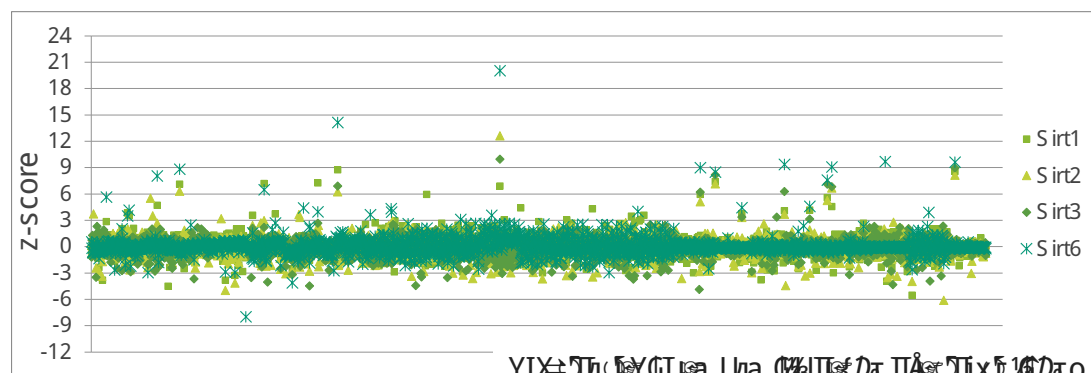
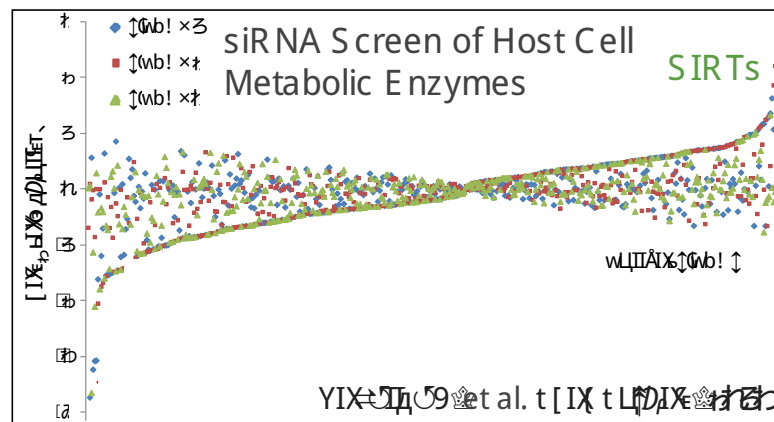
- ǒ Targeting specific a viral protein can lead to drug-resistance
- ǒ One drug generally specific for one virus
- ǒ Novel emergent viruses will likely not be affected by marketed DAAs

- ǒ Host targeted antivirals can likely address these issues

Sirtuins (SIRT) ID'd as affecting HCMV growth & SIRT2 modulator ID'd inhibiting HCMV growth

Tom Shenk lab Princeton:

- HCMV virus infection reprograms host-cell metabolism
- Genetic screen identified SIRT1-7 as viral restriction factors
- A 13 K small molecule screen ID'd 85 SIRT modulators; 61/85 cpds active vs. HCMV in secondary screen



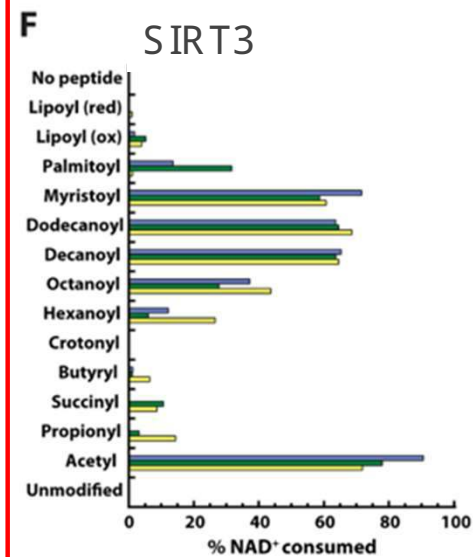
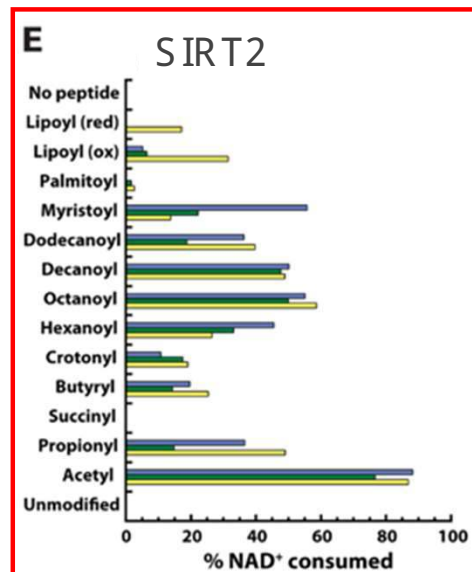
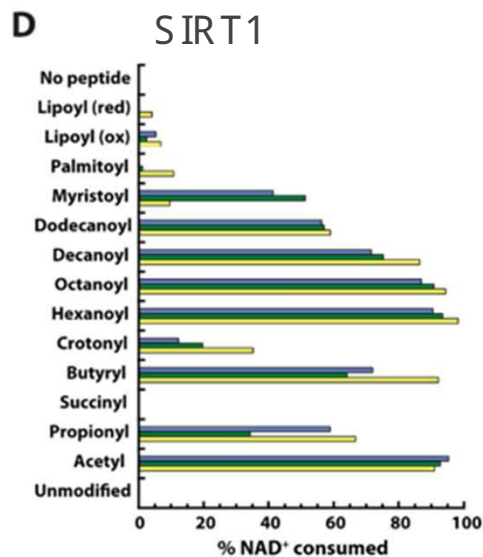
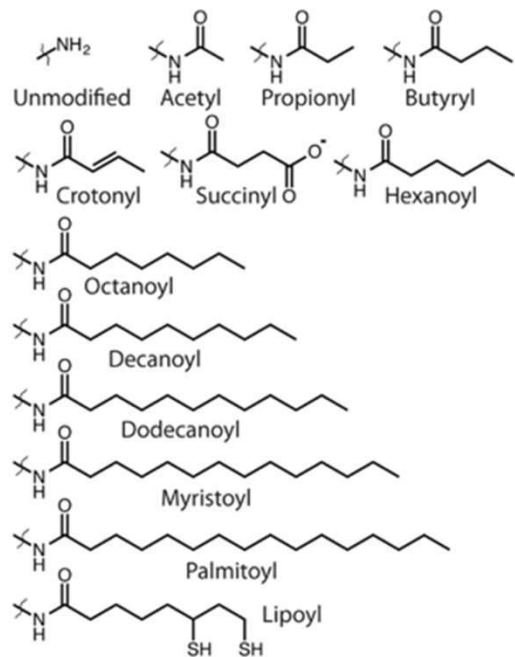
Host Target: Human Sirtuin 2 Protein

- ǒ Sirtuins (SIRTs) are NAD⁺-dependent Class 3 lysine de-acylases (KDACs) that regulate transcription, genome stability, cellular signaling, and energy metabolism in response to the metabolic status of the cell
- ǒ Sirtuin family of 7 proteins vary in sub-cellular location and function to remove a large variety of protein acyl-lysine modifications
- ǒ Viral infection disrupts the metabolic status of the cell and depends on sirtuin regulated functions to replicate productively
- ǒ SIRT2 modulators (nicotinamide, sirtinol, cambinol, AGK2) can engage cellular reprogramming to inhibit cytomegalovirus^{1,2}, hepatitis A virus³, hepatitis B virus^{4,5}, Listeria^{6,7}, Salmonella⁸, Tuberculosis⁹
- ǒ SIRT2 KO mouse is healthy and less susceptible to infection¹⁰
- ǒ Evrys SIRT2 targeted drugs are allosteric enzyme activity modulators

^{1,2}Mao 2016, ³Koyuncu 2014, ^{4,5}Piracha 2018, Yu 2018, ^{6,7}Eskandarian 2013, Pereira 2018, ⁸Gogoi 2018, ⁹Bhaskar 2020, ¹⁰Ciarlo 2017

SIRT1, 2 & 3 are closely related and deacylate various lysine acyl groups

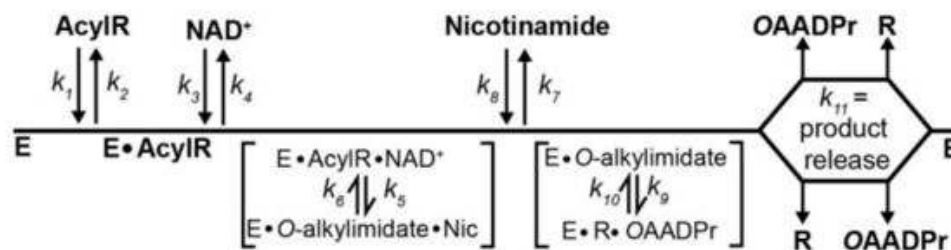
GlnThrAlaArg-Lys(Acyl)-SerThrGlyGly



gfr o cp"LN"Dcg|c"L."FgpwLO 0L"DkqrEj go 04235

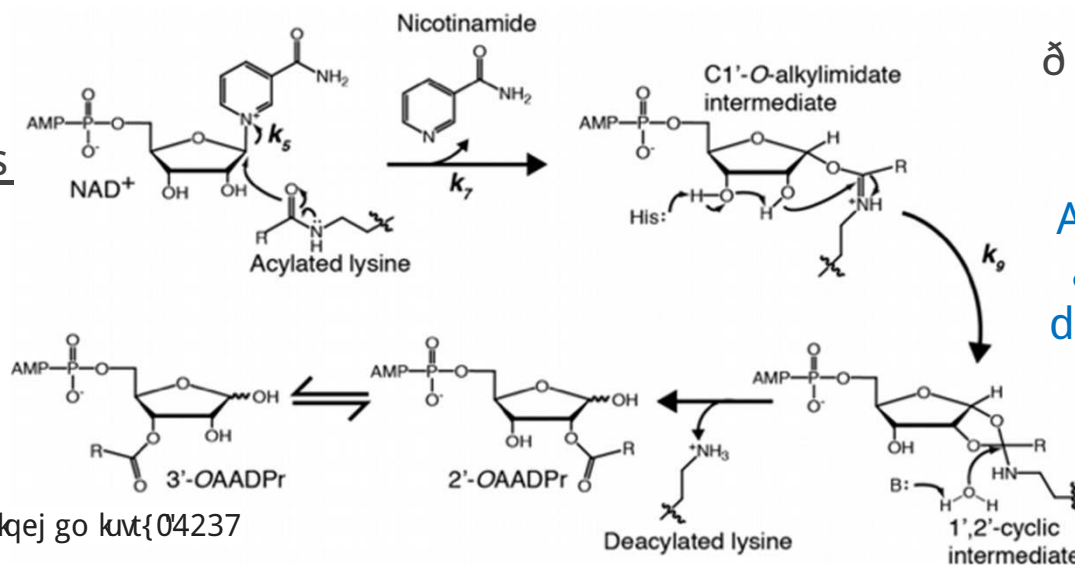
SIRT2 enzyme reaction is complex with multiple substrates, products, transition states, and rate constants

Reaction Mechanism:
 2 substrates
 Ordered binding
 Acyl-peptide, then NAD⁺



Products:
 Nicotinamide
 O-Acyl-ADP-ribose
 de-Acyl-peptide

Chemical Intermediates

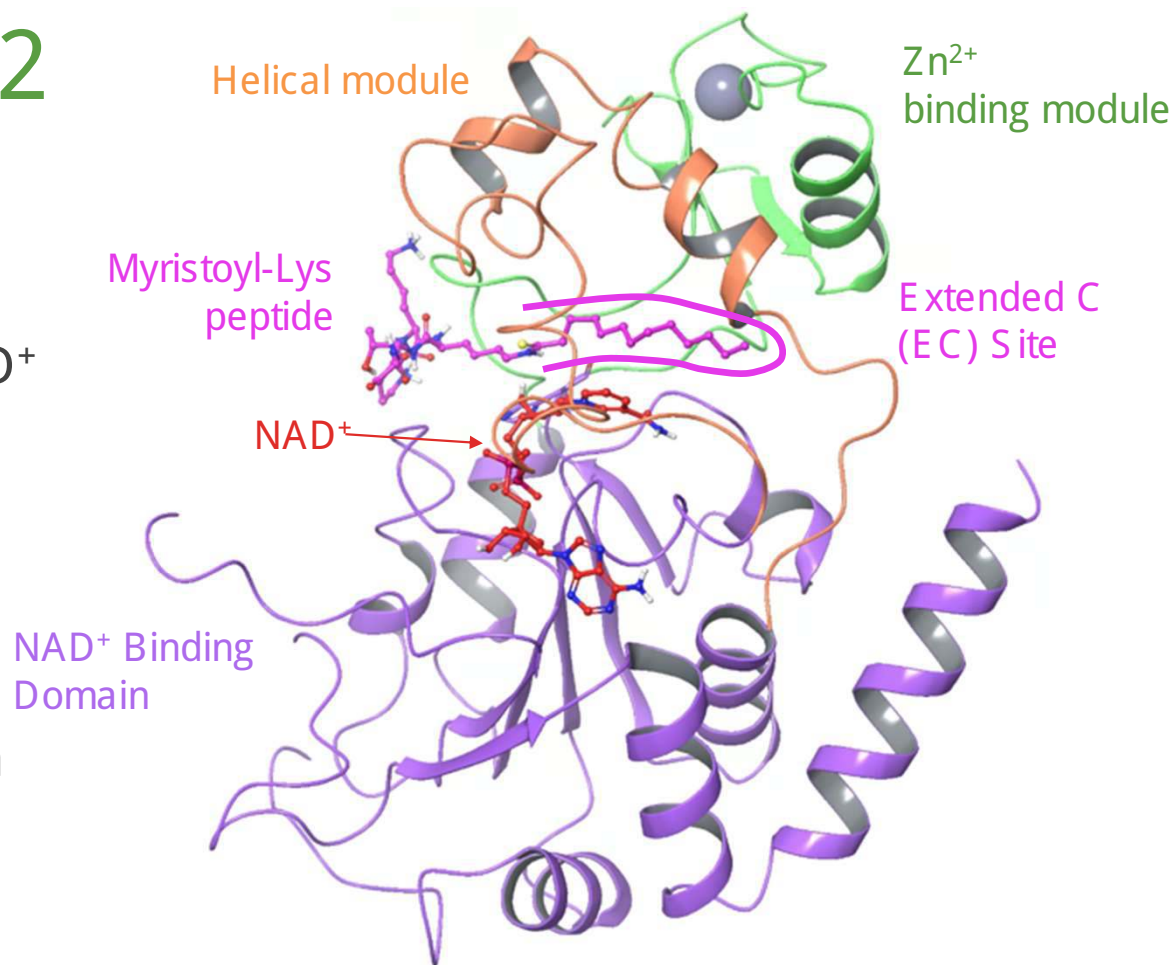


HTMS assay, detecting:
 Ac-ADPR
 De-acyl-peptide
Advantage: Can use any acyl chain length since de-acyl-peptide the same

Hgrf o cp"LN"et al."LO 0Dkqej go kvf{04237

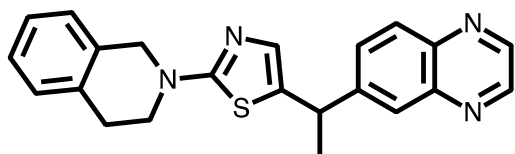
Structure of SIRT2

- SIRT2 X-ray bound to a myristoylated peptide and NAD⁺ (PDB 4X3P)
- The myristoyl binding site (EC site) is induced by the binding of the peptide (or small molecules) and is not present in the apo-protein



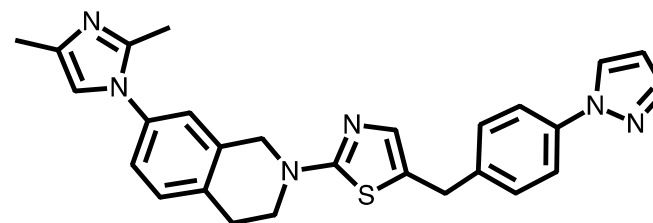
Hit to Lead

• A 13 K small molecule screen at Princeton ID'd FH003 as an antiviral SIRT2 modulator



Princeton: FH003
HCMV IC₅₀ = 8.7 μM
SIRT2 EC₅₀ = 4.1 μM

Evrys
→
Med Chem



Evrys: FLS-359
HCMV IC₅₀ = 0.5 μM
SIRT2 EC₅₀ = 3.0 μM
Wang, Y. et al. W. J. Li et al. W. J. Li et al.

FLS-359: Host targeted SIRT2 modulator results in broad-spectrum antiviral efficacy in vitro

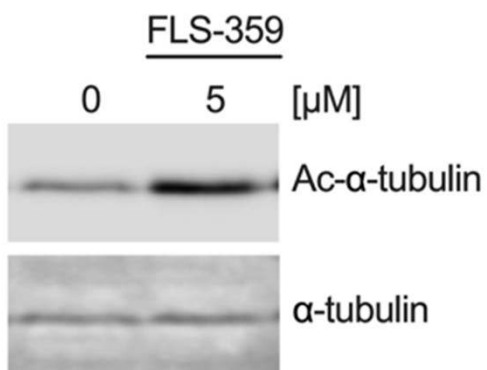
Virus/ Host Cell	Virus Family	FLS-359 IC ₅₀ (I M)	Host Cell CC ₅₀ (I M)	¹ SI CC ₅₀ /IC ₅₀	² SOC/C IC ₅₀ (I M)	SOC/ Comparator (C)
³ SARS-CoV-2/Human Calu3	ϕ-Coronavirus	0.3	15.8	52.7	0.4	Remdesivir (C)
³ Zika/Human HFF	Flavivirus	0.4	41.6	104.0	2.8	Amodiaquine (C)
⁴ HC MV/Human MRC5	ϕ-Herpesvirus	0.5	>15.8	>40	2.7	Ganciclovir (SOC)
⁵ Influenza A/Human dNHBE	Orthomyxovirus	⁷ 1.2	>100	>83.3	⁷ 0.7	Ribavirin (C)
⁴ HC oV-OC 43/MRC5	ϕ-Coronavirus	1.7	>50	>30.1	0.1	Remdesivir (C)
³ J unin/Human HFF	Arenavirus	3.2	>25	>7.8	0.2	RIID E-1 (C)
⁶ Hepatitis B/Human PHH	Hepadnavirus	4.8	>10	>2.1	0.03	Tenofovir (SOC)
Epstein-Barr/Human Akata	γ-Herpesvirus	3.8	>100	>26.3	43	Ganciclovir (C)
RSV/Human MRC5	Orthopneumovirus	6.7	>12.5	>1.9	16.1	Ribavirin (SOC)

Results are shown for representative assays. ¹SI = selectivity index, ²SOC = standard of care or C = comparator compound, ³HFF = human foreskin fibroblasts, ⁵dNHBE = differentiated normal human bronchial epithelial cells, PHH = primary human hepatocytes, ⁷IC₉₀.

Wang Y et al. WJG 2019; 33(11): 2111-2120

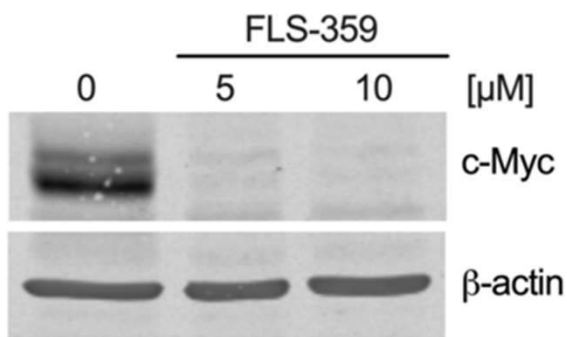
FLS-359 affects known cellular targets of SIRT2

From HepG2 cells

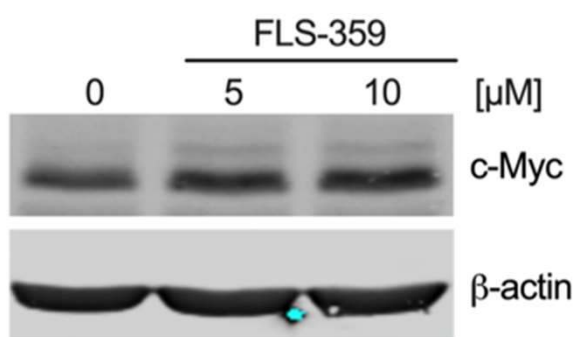


HepG2 cells were treated with FLS-359 plus 1 μ M trichostatin A for 24 h and levels of acetylated a-tubulin (Ac-a-tubulin) were determined by Western blot. Total a-tubulin was monitored as a control

From MDA-MB-231 cells



From MRC-5 cells



MDA-MB-231 cells or MRC-5 fibroblasts were treated with drug for 72h and c-Myc was monitored by Western blot with b-actin as a control.

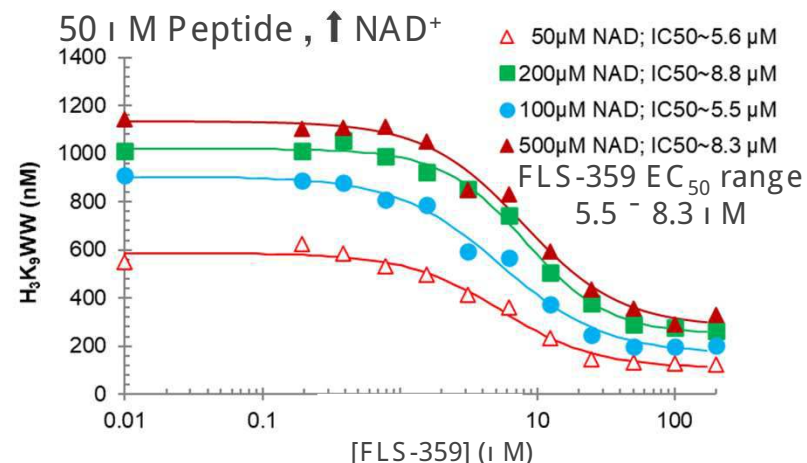
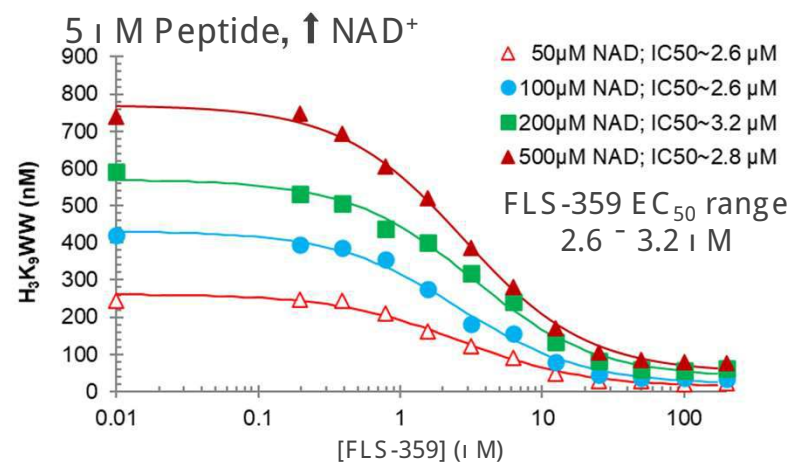
Wang et al. WJG 2011;23(12):2111-2116

Evrys FLS-359 SIRT2 modulator is not a classic competitive inhibitor

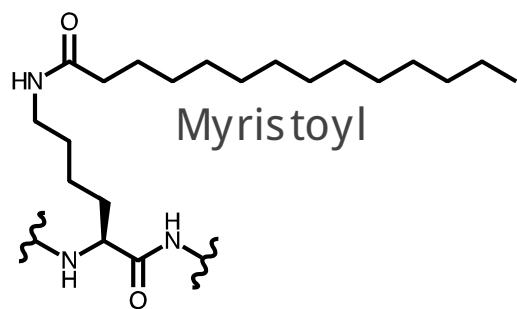
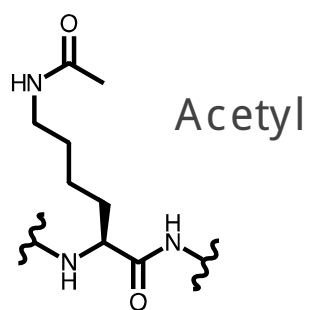
- ∅ SIRT2 Acetyl-peptide hydrolysis rate is reduced
- ∅ Effect not altered when raising [NAD]
- ∅ Effect marginally altered raising [Ac-peptide]
- ∅ Cpds not fully competitive with substrates
- ∅ SIRT2 activity is not reduced to 0
- ∅ Compounds are SIRT2 modulators
- ∅ Evrys compounds are selective for SIRT2

Drug	SIRT1 de-Acetyl EC ₅₀	SIRT2 de-Acetyl EC ₅₀	SIRT3 de-Acetyl EC ₅₀
FLS-359	>100 μM	3 μM	>100 μM

Wang Y et al. WJPT 2023



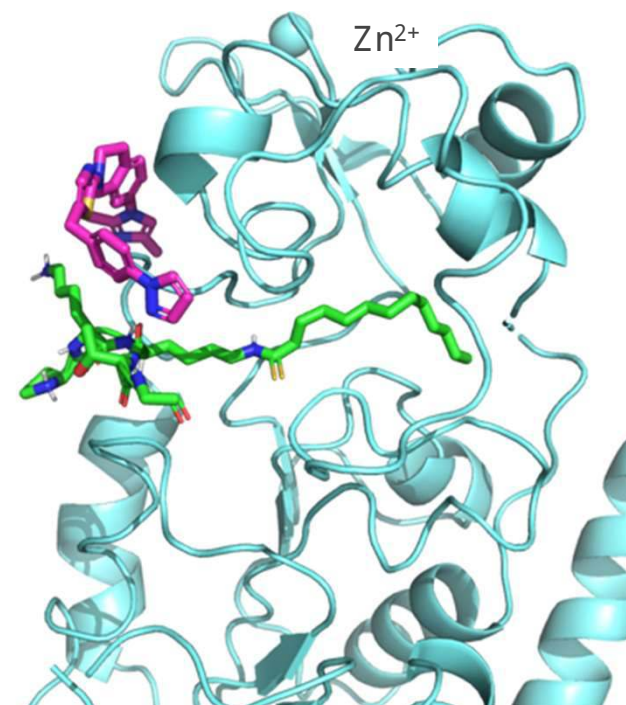
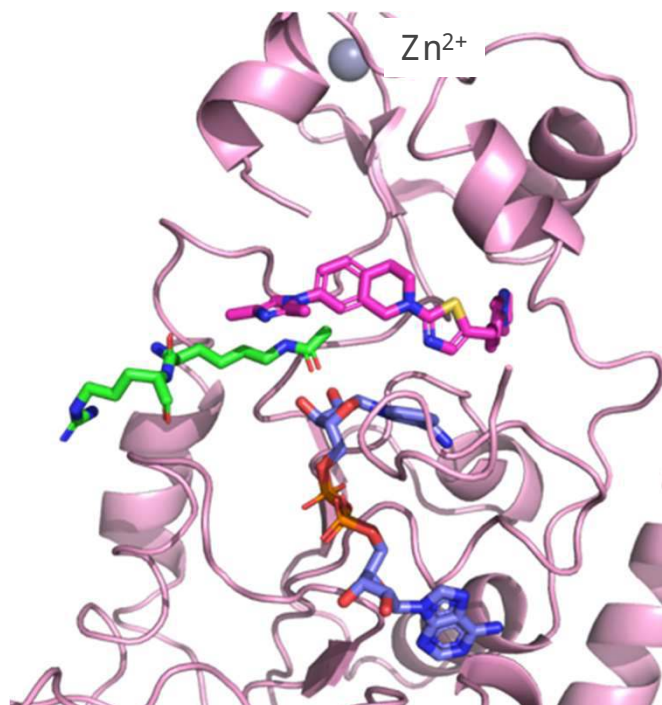
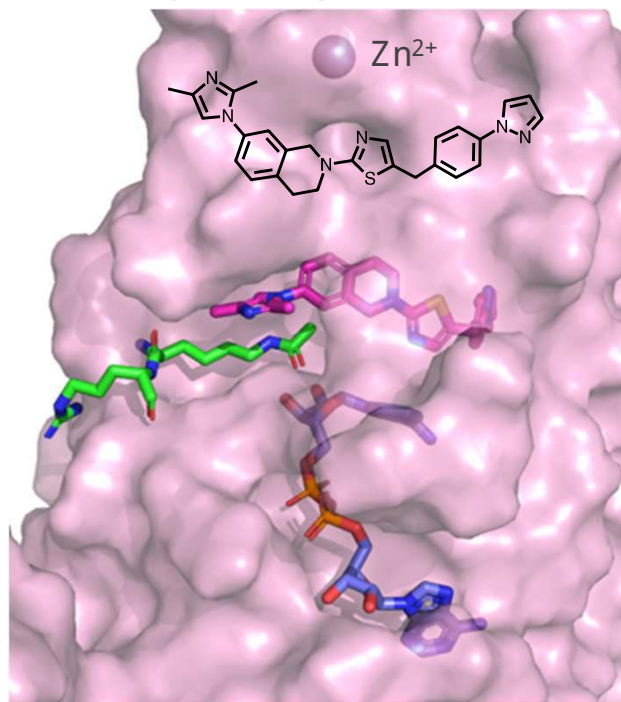
FLS-359 is a selective peptide substrate modulator



Cpd	Structure	Deacetylase IC ₅₀ (I M)	Demyristoylase IC ₅₀ (I M)
FLS-359		3.0	>100
SirReal2		0.2	>100
AGK2		0.5	>100

Wang et al. WJG 2011;23(11):1911-1918

SIRT2 can accommodate FLS-359, NAD⁺, and Acetyl- but not myristoylated-substrate

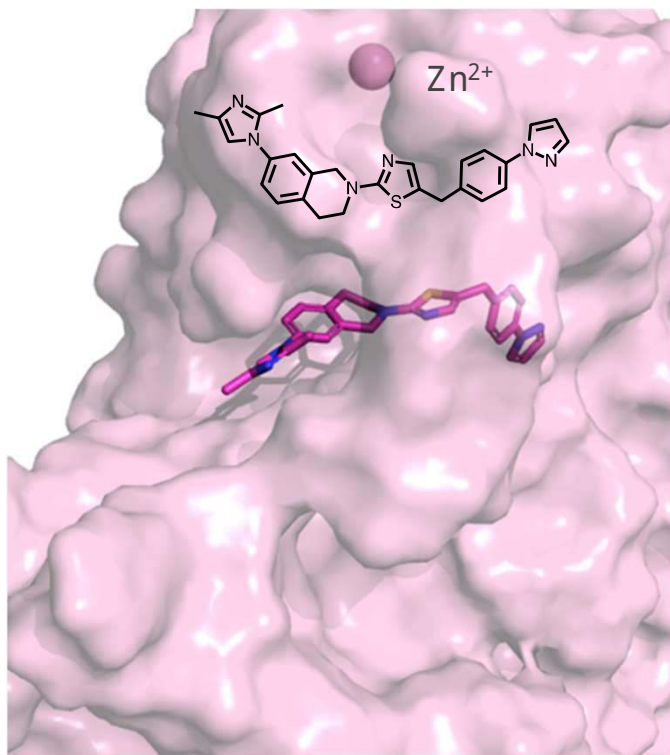


FLS-359 docked to SIRT2 with acetylated lysine PDB 4RMH. Binding of FLS-359 (magenta sticks) does not prevent binding of acetyl lysine substrate (green sticks) or NAD⁺ (purple sticks, from PDB 4RMG). Left panel: SIRT2 with surface.

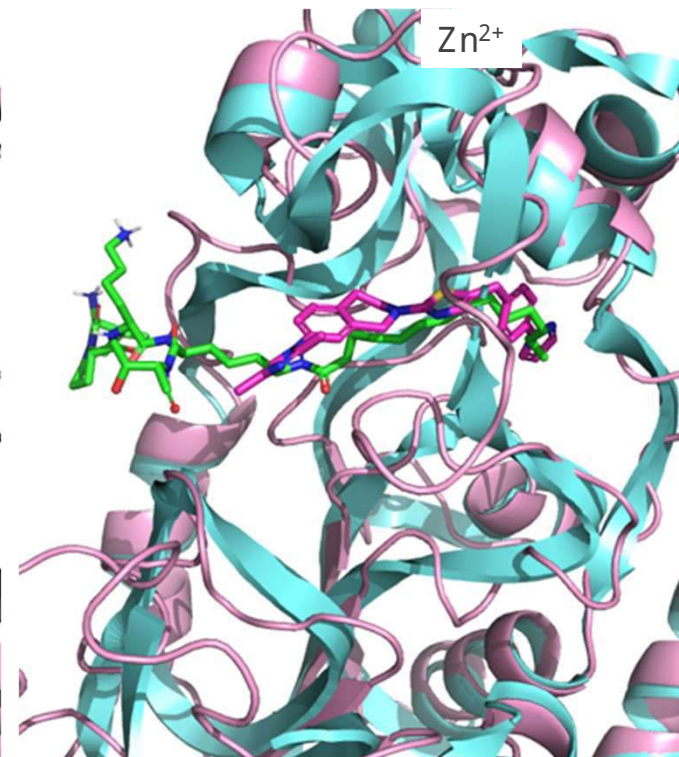
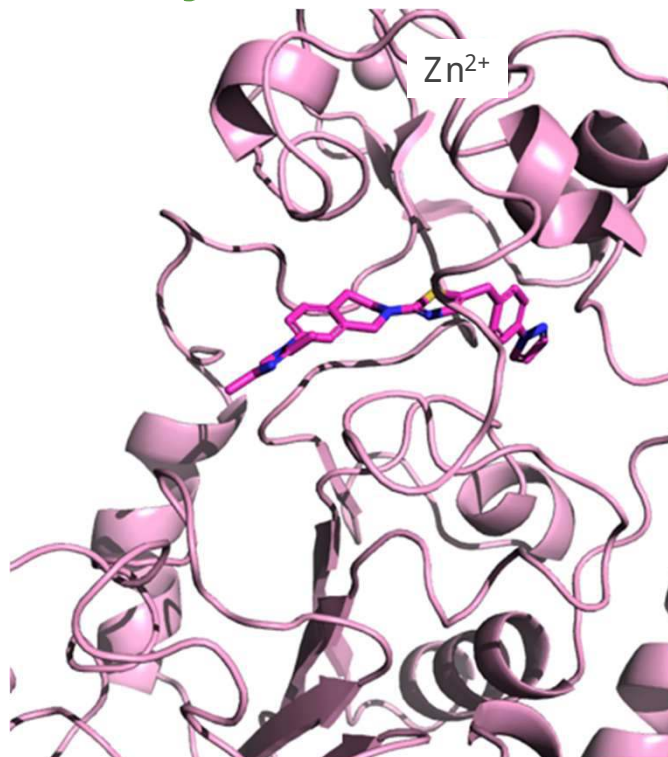
Center panel: SIRT2 with ribbon structure only. Xie, Y. et al. WQJLJ 2019

Myristoylated peptide (green sticks, PDB 4Y60) competes for binding with FLS-359 (magenta sticks) preventing compound binding to SIRT2.

FLS-359⁻ SIRT2 co-crystal PDB 7T1D



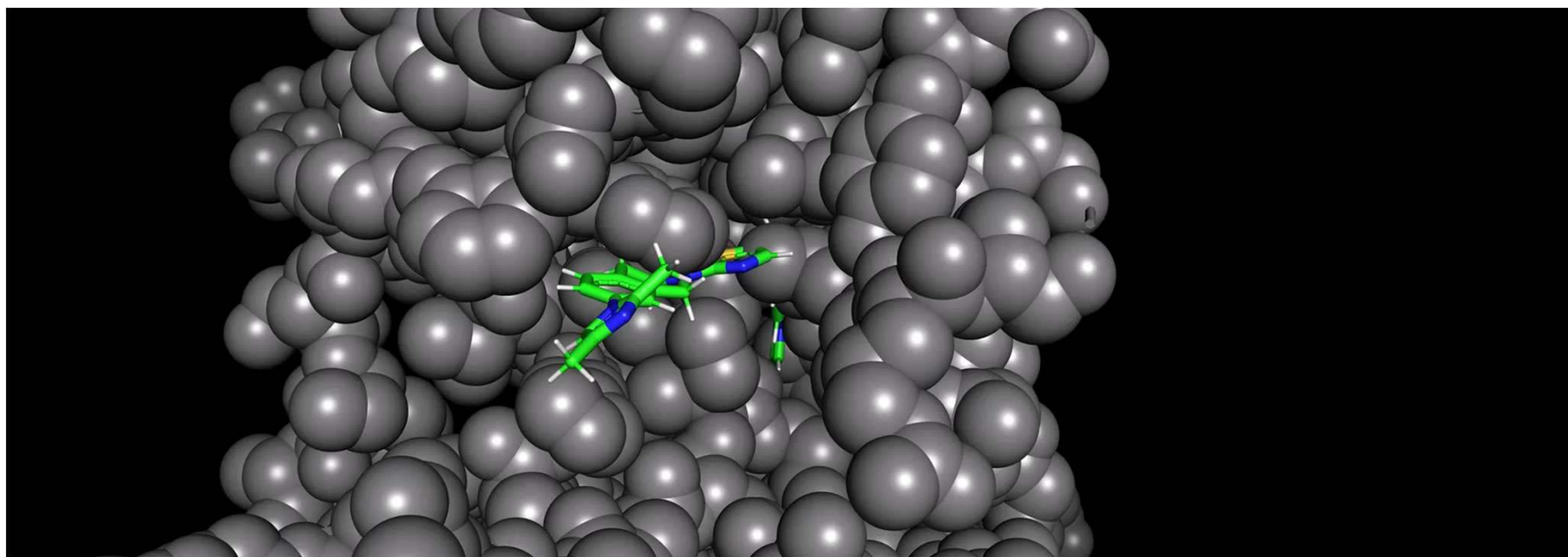
FLS-359 SIRT2 cocystal shows compound filling EC site.
Roche KL, et al. J Clin Invest. 2023



FLS-359 SIRT2 cocystal 7T1D (pink)
overlayed with myristoylated
peptide 4Y60 (cyan).

EC site is induced by FLS-359 binding to SIRT2

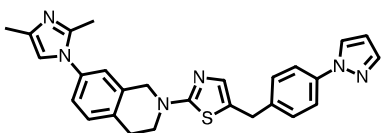
Start: Apo-SIRT2 X-ray; End: FLS-359-bound SIRT2 X-ray, conformation change



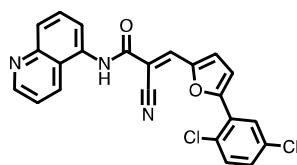
Published SIRT2 modulators reduce HCMV yield in vitro

Compound	HCMV IC ₅₀ (I M)	MRC-5 CC ₅₀ (I M)	SI: CC ₅₀ /IC ₅₀	SIRT2 Deacetyl IC ₅₀ (I M)
FLS-359	0.5	>15.8	>40	3.0
AGK2	3.4	>100	>29.4	3.5
AK-7	8.2	>25	>3	15.5
MIND4	14.4	19.8	1.4	3.5
SirReal2	14.9	>20	>1.3	0.2
TM	26.0	46.3	1.8	0.03

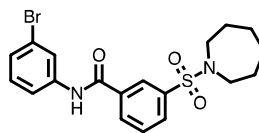
FLS-359



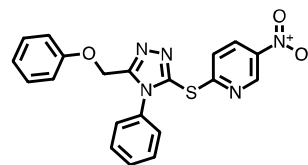
AGK2



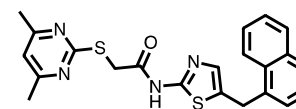
AK7



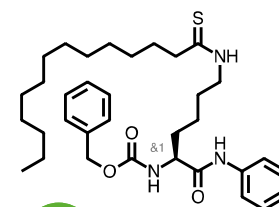
MIND4



SirReal2

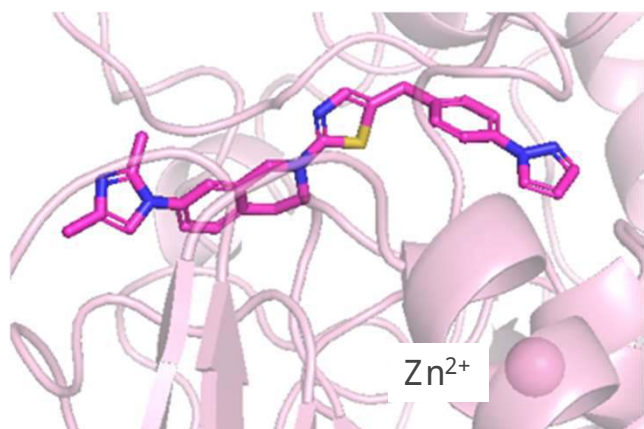


TM

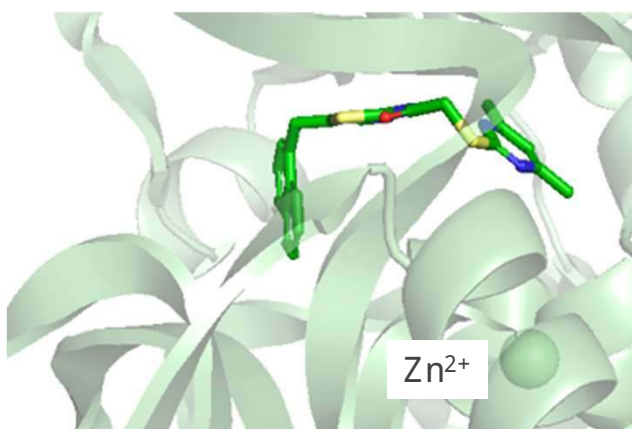
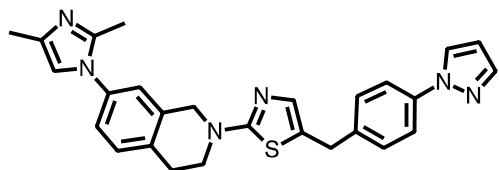


Wang et al. WJPT 2019

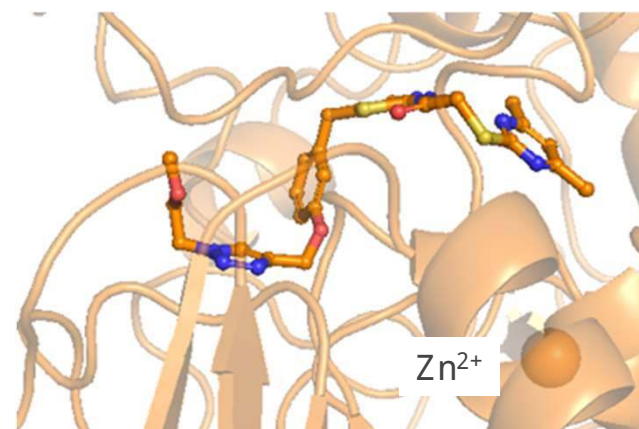
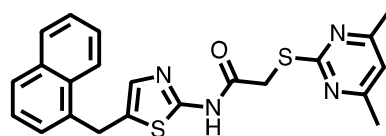
Small molecule SIRT2 modulators have different SIRT binding conformations



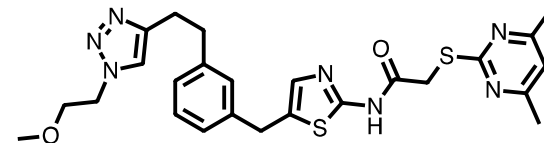
FLS-359 7T1D



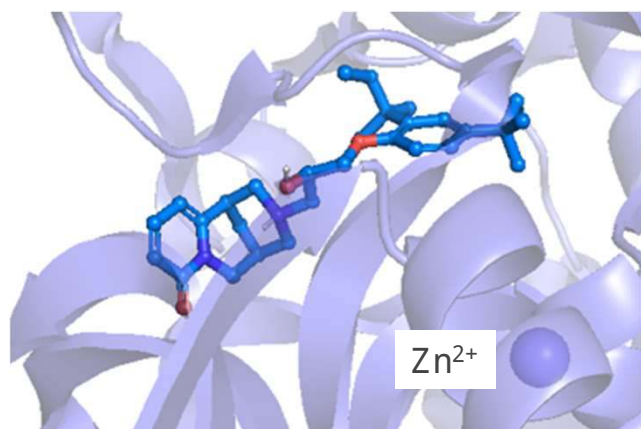
SirReal2 4RMG



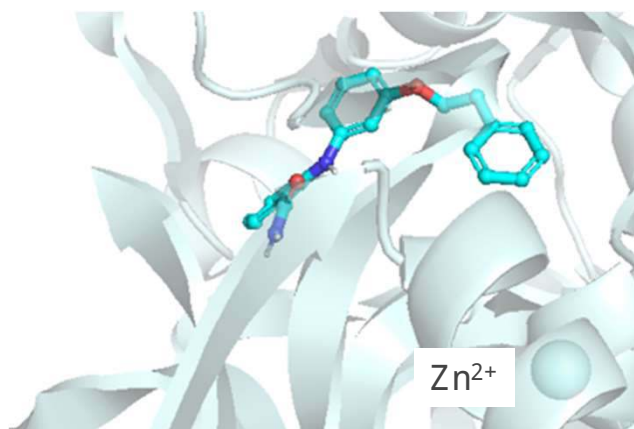
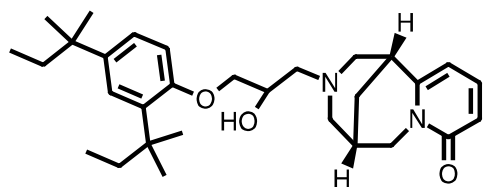
SirReal1.2 8OWZ



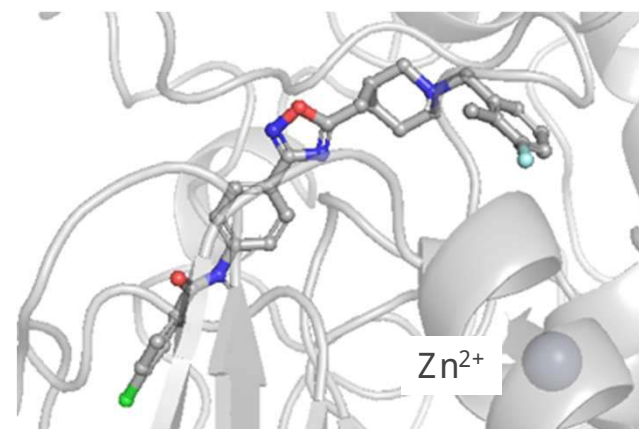
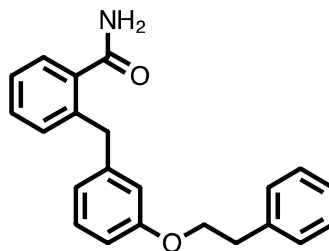
Small molecule SIRT2 modulators have different SIRT binding conformations



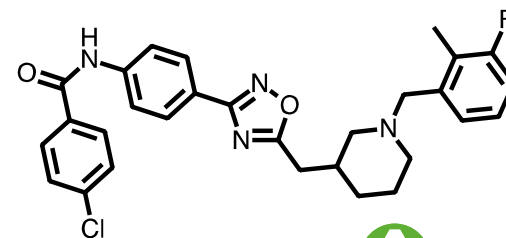
NPD11033 5Y0Z



Benzamide 6 5Y5N



Oxadiazole 16 8PY3



Summary

- An unbiased siRNA genetic screen in HCMV infected fibroblasts identified the enzyme class Sirtuins as viral restriction factors
- A small molecule SIRT enzyme screen found 85 SIRT modulators with 65 having anti-HCMV activity
- SIRT2 modulation affects multiple human pathogens
- SIRT2 can deacylate lysines with chain lengths ranging from C2 acetyl to C14 myristoyl
- The mechanism of SIRT2-mediated deacylation is multistep

Summary

- A variety of SIRT2 modulators affect HCMV replication
- Evrys lead optimization resulted in FLS-359, which allosterically modulates SIRT2
- Host-targeted FLS-359 has broad-spectrum antiviral activity
- The EC pocket is absent in apo-SIRT2 and is induced by substrate and small molecule binding
- The EC pocket is flexible and can accommodate a variety of small molecule binders that adopt very different conformations in their interactions with SIRT2

Acknowledgements

Evrys Bio

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