

Chemical Lysine Modification at a single site

2018/12/20

M1 Murata

Contents

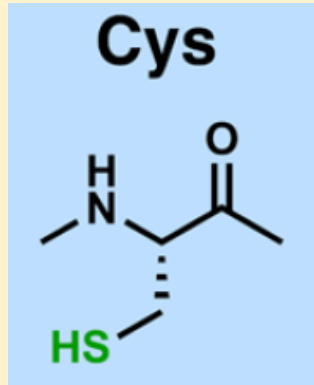
- Amino acid target for bioconjugation
- Chemical Lysine modification
 - *N*-hydroxysuccinimide (NHS)-ester
 - α , β -unsaturated sulfonamide
 - Sulfotetrafluorophenyl esters
 - Stilbene
 - β -lactam
 - Sulfonyl acrylate (Lysine activated)
- Summary

Contents

- Amino acid target for bioconjugation
- Chemical Lysine modification
 - *N*-hydroxysuccinimide (NHS)-ester
 - α,β -unsaturated sulfonamide
 - Sulfotetrafluorophenyl esters
 - Stilbene
 - β -lactam
 - Sulfonyl acrylate (Lysine activated)
- Summary

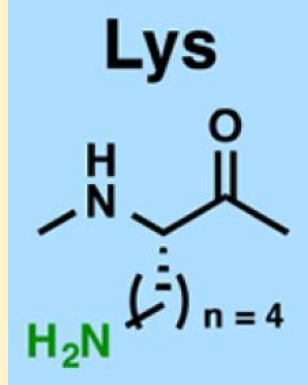
Amino acids for bioconjugation

Conventional



(pKa=8.7)

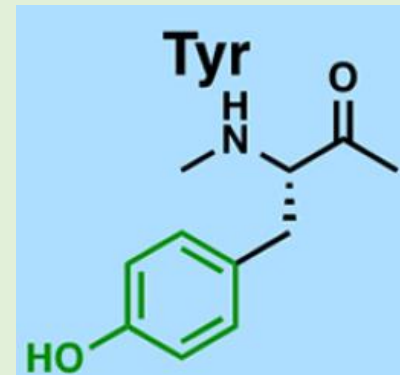
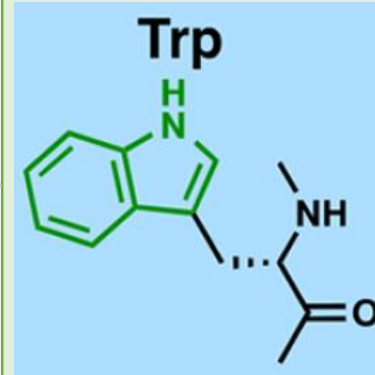
- ✓ Reactive because of nucleophilicity
- ✓ Abundant (Lys)



(pKa=10.5)

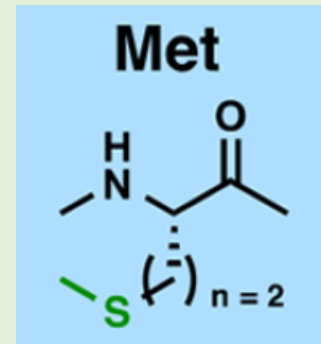


Recent



(pKa=10.46)

- ✓ Less-reactive
- ✓ Rare



Trp... keto-ABNO

Tyr... reactivity can be efficiently tuned by **pH control**

Met...only native residue that can be alkylated under **acidic conditions**

Cys and Lys residues for bioconjugation

The advantage of targeting Cys or Lys residues

- ① abundant (Lys)
- ② Easy to target because of high reactivity

	Ratio in human protein(%)	The position of residue	Comments
Cys	1.9	within the folded proteins	Not accessible
Lys	5.9	On the surface of proteins	accessible

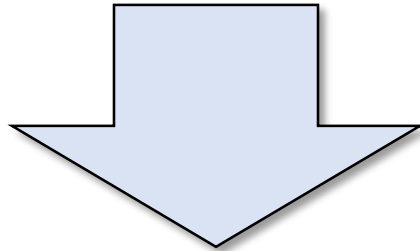
Native lysine residues are more convenient targets for protein modification than cysteine residues,

Targeting naturally occurring side chains in a chemo- and region-selective fashion remains a great and unexplored challenge.

Problems in Lys-bioconjugation

Problem

- Conventional method –
 - ✓ Difficulties to target specific single Lys residue
 - ✓ Require a large excess of often-valuable reagents. (e.g. 10 eq)
 - ✓ Require metals (e.g. Ru, Pd...) or other additives that must later be removed from the reaction mixture.

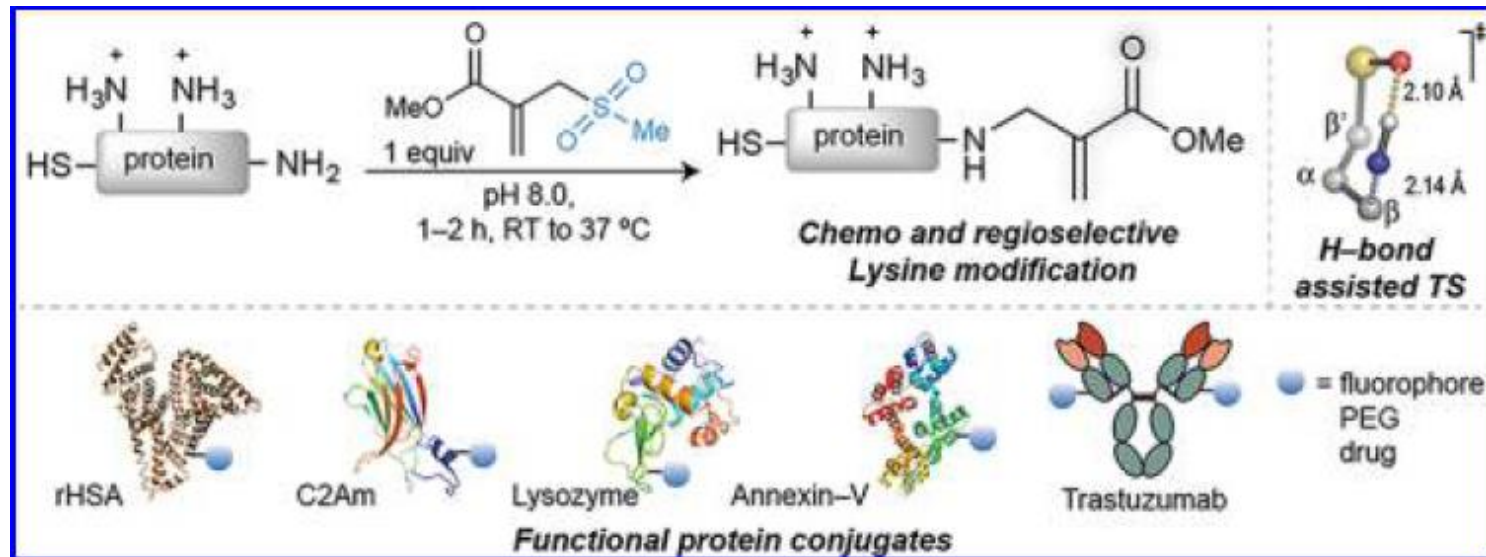


“A general way to selectively target single Lysine is lacking”

A main literature discussed in this seminar

- Sulfonyl acrylate (Lysine activated) -

Hydrogen bond assisted chemo- and regioselective modification of lysine on native proteins



- ✓ Stoichiometric amount of sulfonyl acrylate reagent
- ✓ Proceed to completion rapidly (1~2h)
- ✓ Under mild conditions (pH=8.0, rt-37°C)
- ✓ Applicable to a range of native protein types
- ✓ The products are stable
- ✓ The method is compatible with Cys bioconjugation method

Contents

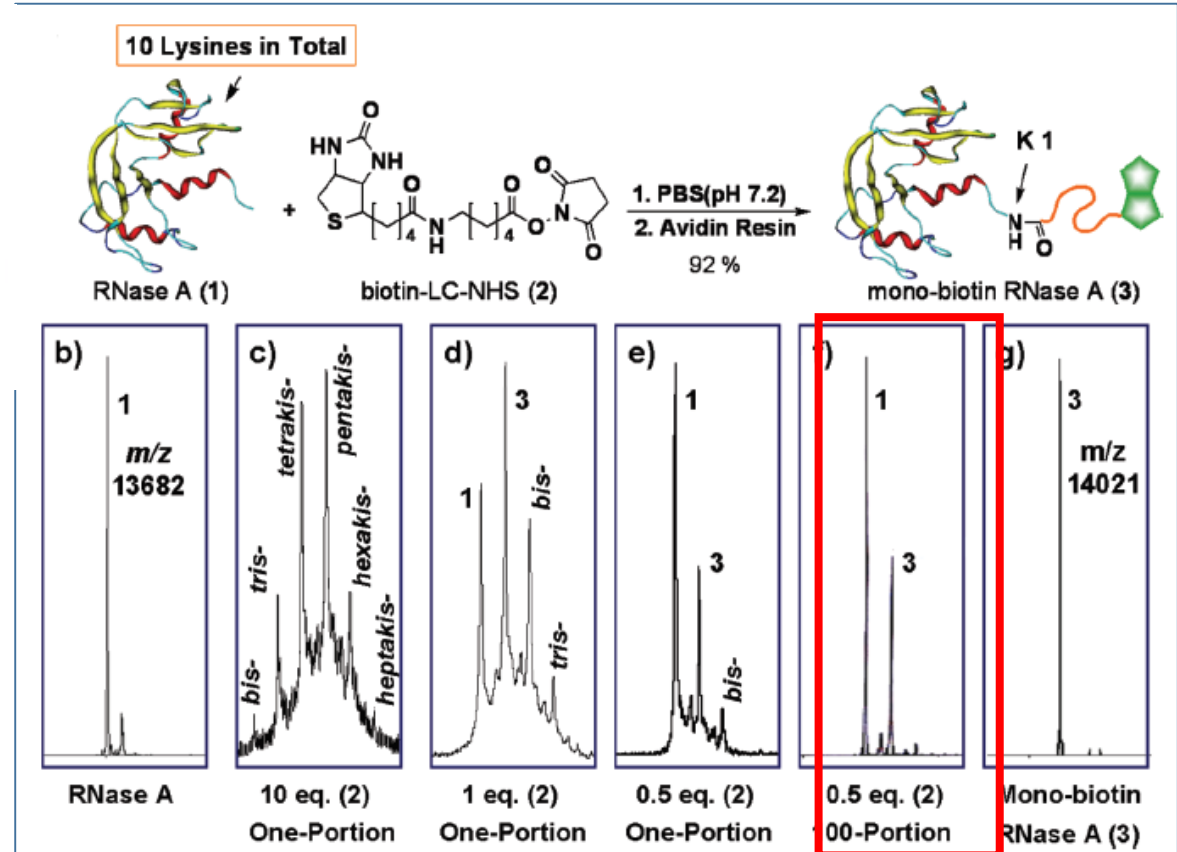
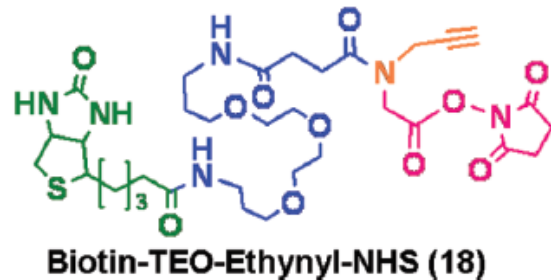
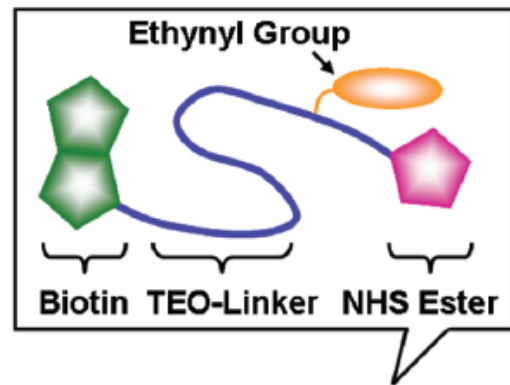
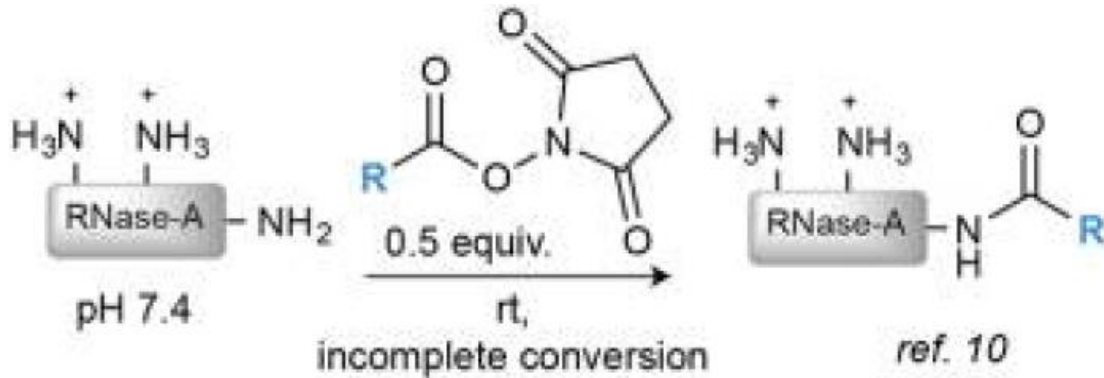
- Amino acid target for bioconjugation
- **Chemical Lysine modification**
 - *N*-hydroxysuccinimide (NHS)-ester
 - α,β -unsaturated sulfonamide
 - Sulfotetrafluorophenyl esters
 - stilbene
 - β -lactam
 - Sulfonyl acrylate (Lysine activated)
- Summary

Chemical Lysine modification

- *N*-hydroxysuccinimide (NHS)-ester
- α,β -unsaturated sulfonamide
- Sulfotetrafluorophenyl esters
- Stilbene
- β -lactam

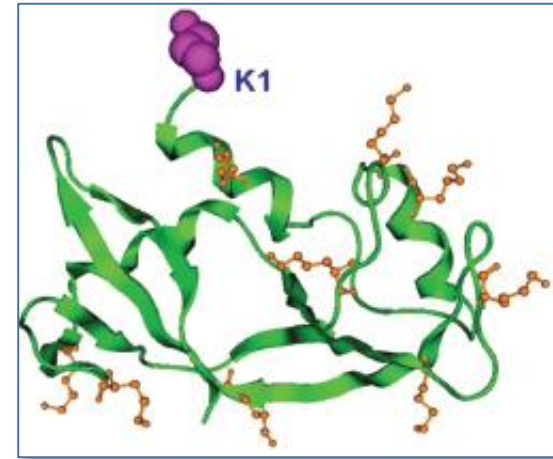
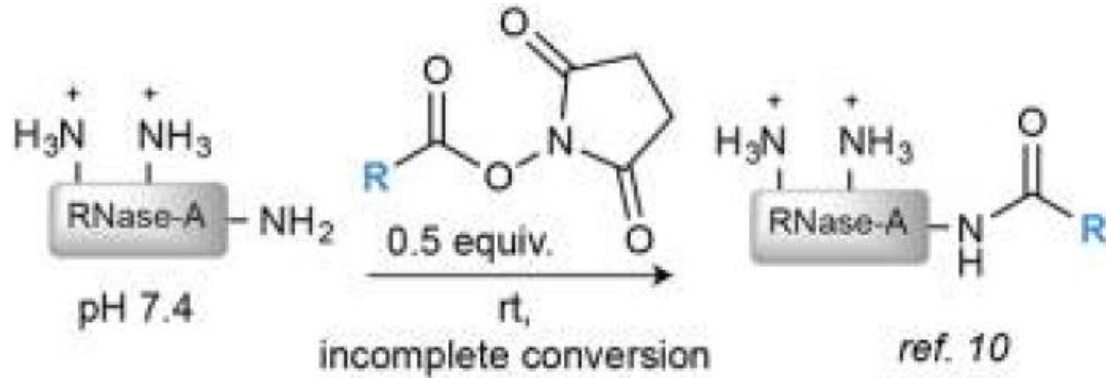
Chemical Lysine modification 1

- N-hydroxysuccinimide (NHS)-ester type -



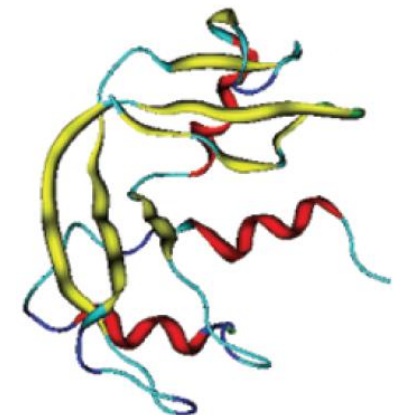
Chemical Lysine modification 1

- N-hydroxysuccinimide (NHS)-ester type -

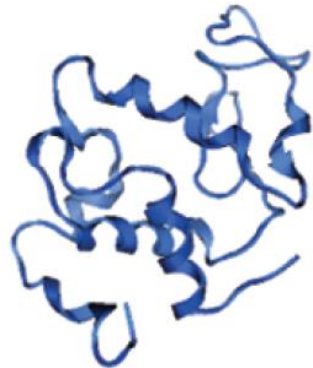


Rnase A

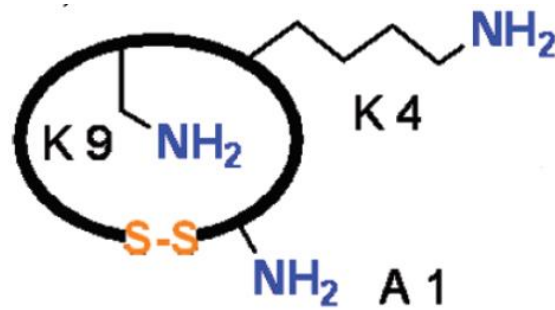
✓ the highest solvent accessibility.



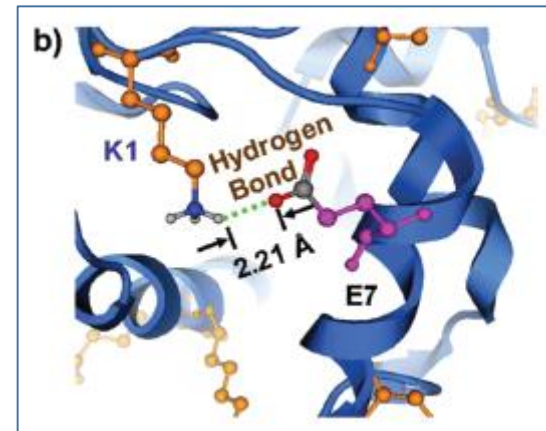
RNase A (1)



Lysozyme C



SST-14

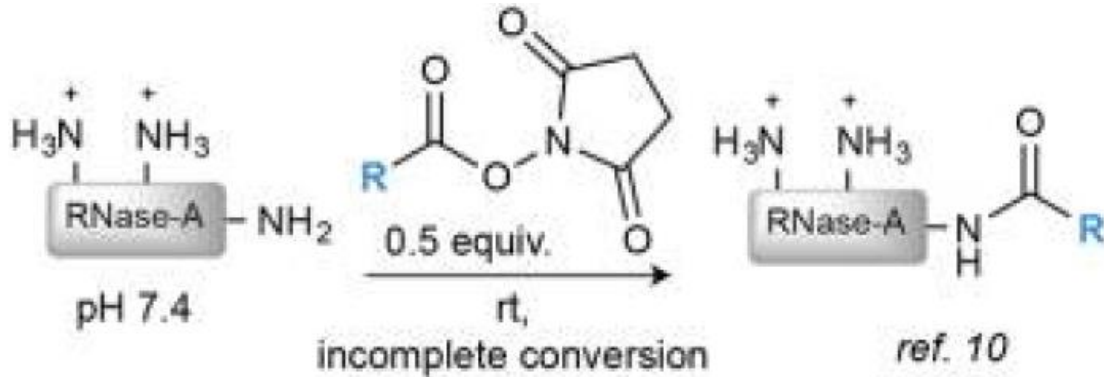


Lysozyme C

✓ hydrogen bond interaction between K1 and E7

Chemical Lysine modification 1

- N-hydroxysuccinimide (NHS)-ester type -

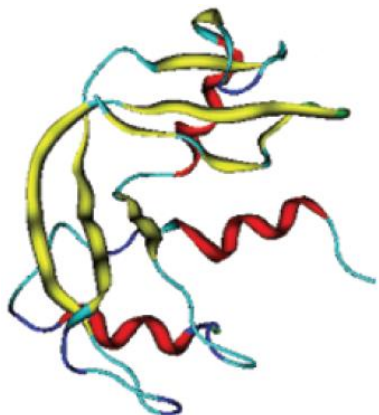


Advantage

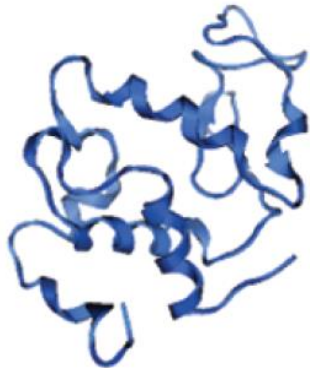
- ✓ Fast (2h)
- ✓ Low cost
- ✓ The risk of misfolding is low

Disadvantage

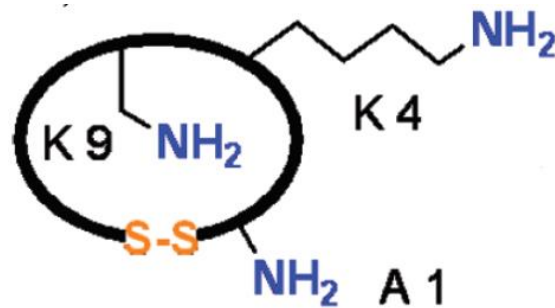
- ✓ the requirement of substoichiometric amounts of reagent
➔ incomplete conversion



RNase A (1)



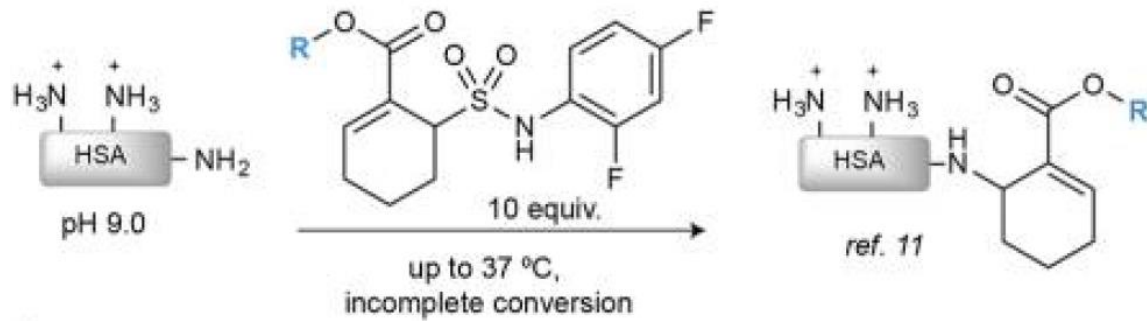
Lysozyme C



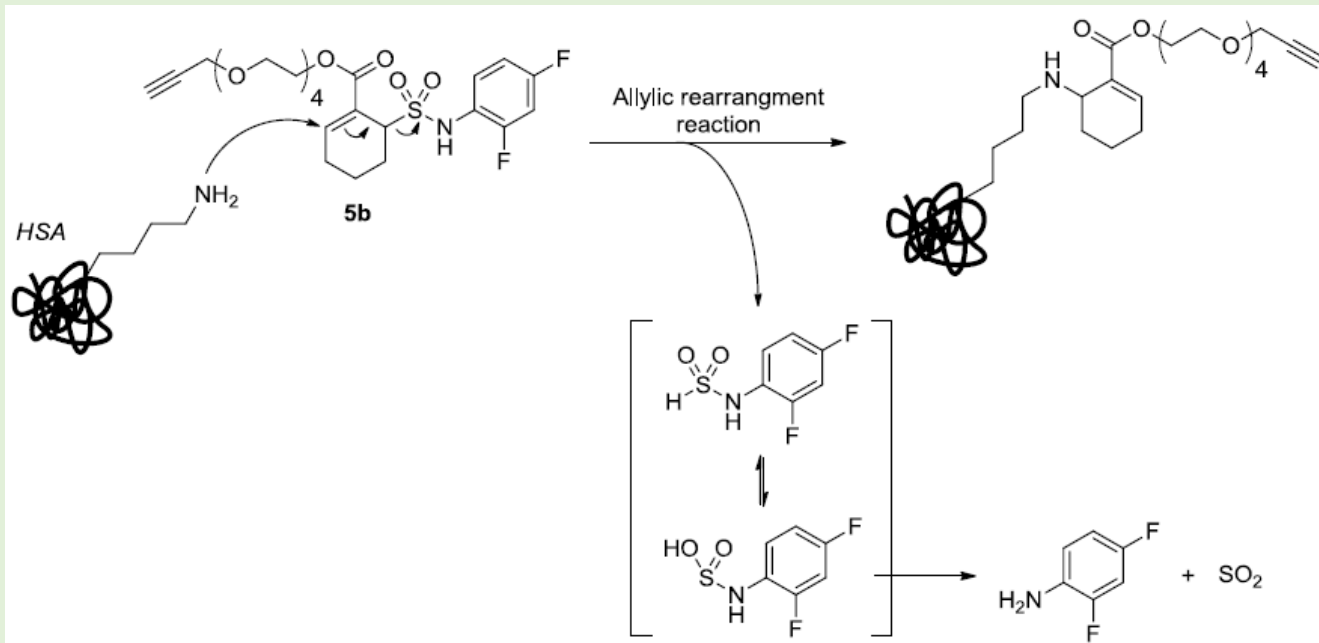
SST-14

Chemical Lysine modification 2

- α,β -unsaturated sulfonamide type -



Proposed Mechanism of the Conjugation Reaction with HSA



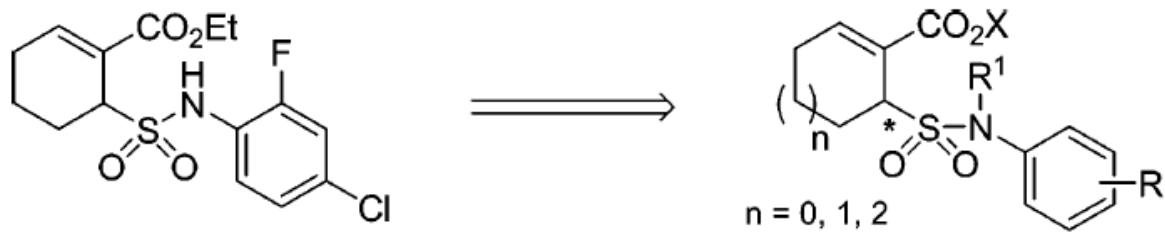
Angew. Chem. Int. Ed. 2014, 53, 11783–11786

Bioconjugate Chem. 2016, 27, 2271–2275

Chemical Lysine modification 2

- α,β -unsaturated sulfonamide type -

Screening of the Takeda chemical library was carried out
To develop a new therapeutic agent for sepsis.



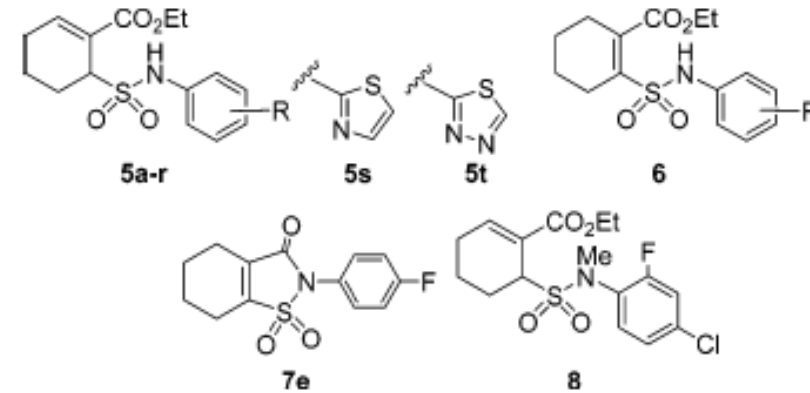
Lead compound

Yamada *et al.*, *J. Med. Chem.* **2005**, *48*, 7457-7467

Barbas *et al.*, *Angew. Chem. Int. Ed.* **2014**, *53*, 11783–11786

Barbas *et al.*, *Bioconjugate Chem.* **2016**, *27*, 2271–2275

Table 1. Inhibitory Activities of *N*-Arylsulfamoyl Derivatives **5a–t**, **6a,b,e**, **7e**, and **8** on NO Production

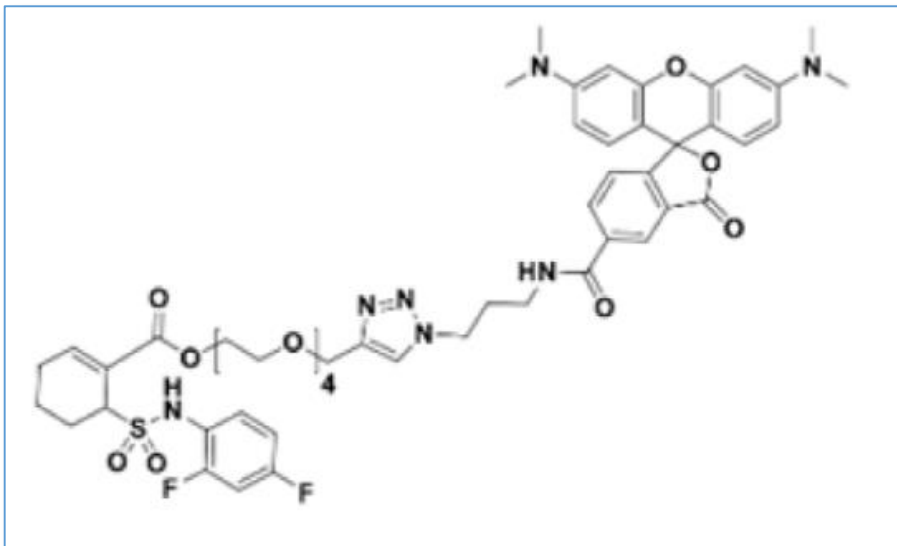
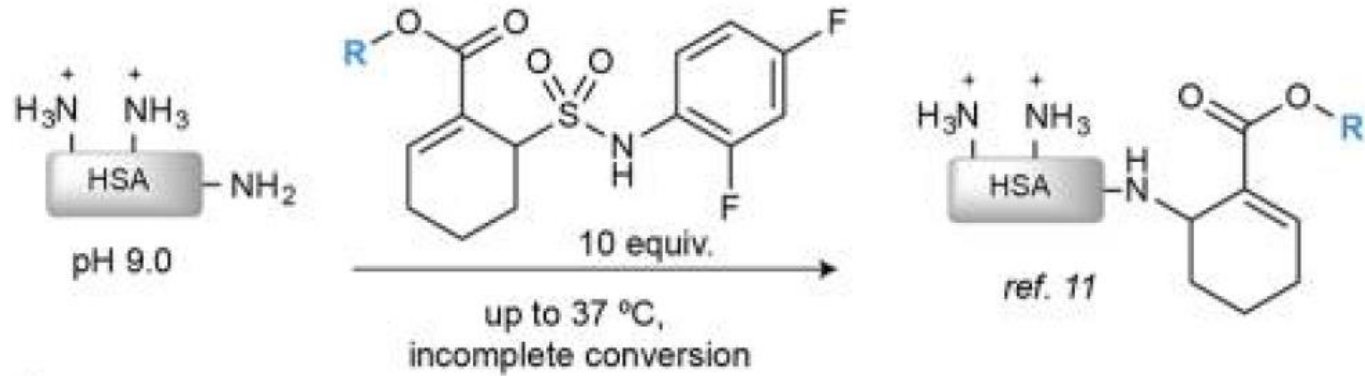


compd	R	IC ₅₀ ^a (nM)	compd	R	IC ₅₀ ^a (nM)
5a	2-F, 4-Cl	160 ± 65	5n	2-Cl, 4-F	3.2 ± 0.89
5b	H	260 ± 113	5o	2-Cl, 4-Me	41 ± 14
5c	2-F	75 ± 45	5p	2-Cl, 4-CN	1600 ± 172
5d	3-F	150 ± 14	5q	2-Et	130 ± 16
5e	4-F	110 ± 11	5r	2-CO ₂ Me	1100 ± 194
5f	2-Cl	12 ± 1.2	5s	–	> 10000
5g	3-Cl	66 ± 19	5t	–	> 10000
5h	4-Cl	400 ± 279	6a	2-F, 4-Cl	1700 ± 495
5i	2,3-F ₂	140 ± 49	6b	H	> 8200
5j	2,6-F ₂	160 ± 4.9	6e	4-F	1400 ± 363
5k	2,4-F ₂	16 ± 1.4	7e	–	4100 ± 1742
5l	2,4,5-F ₃	30 ± 1.8	8	–	230 ± 45
5m	2,4-Cl ₂	20 ± 2.0			

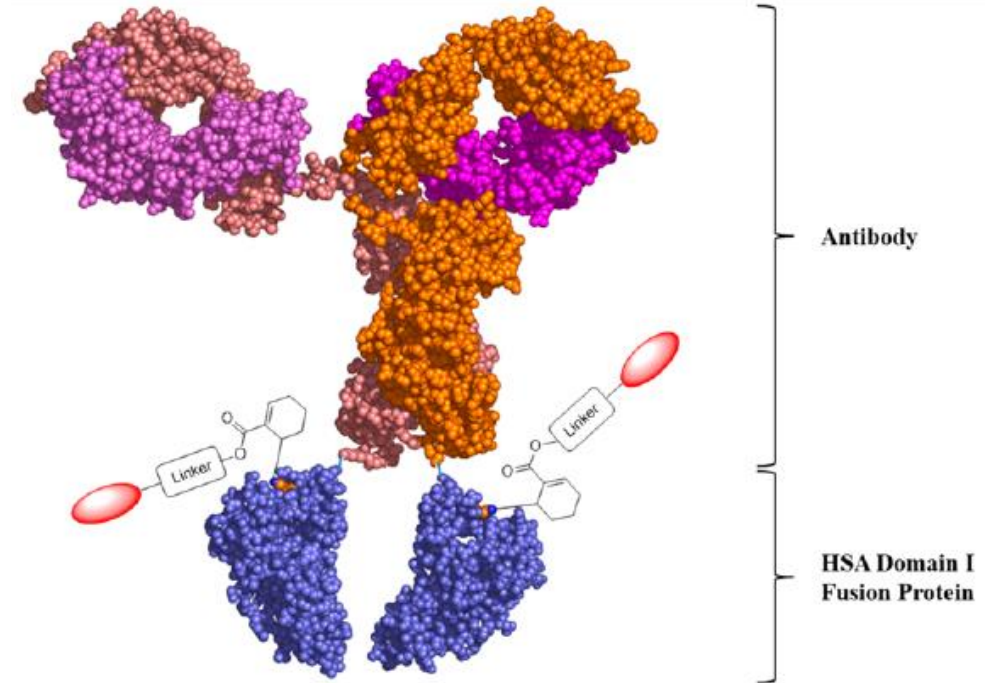
^a The inhibitory activity is shown as an IC₅₀ value, which is the concentration of test compound required to suppress the production of NO by 50% of control. Values are the mean ± SD of two or three experiments.

Chemical Lysine modification 2

- α,β -unsaturated sulfonamide type -



Rhodamine-linked cyclohexene sulfonamide compound (cHx-Rho).



- ✓ rapidly modify HSA
- ✓ excellent serum stability

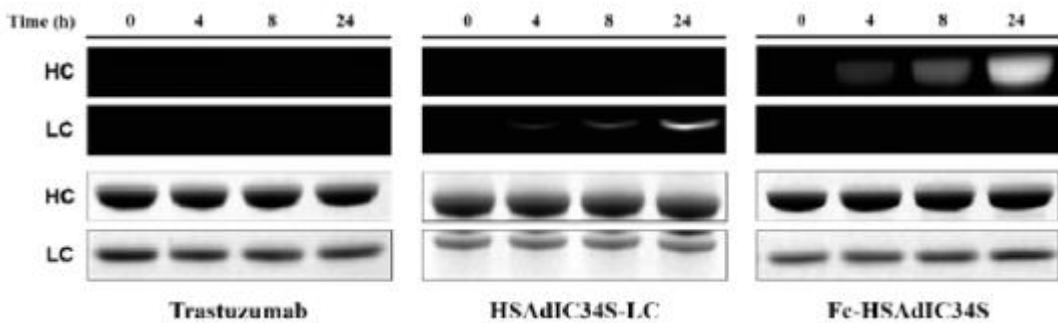
Barbas *et al.*, *Angew. Chem. Int. Ed.* **2014**, 53, 11783–11786

Barbas *et al.*, *Bioconjugate Chem.* **2016**, 27, 2271–2275

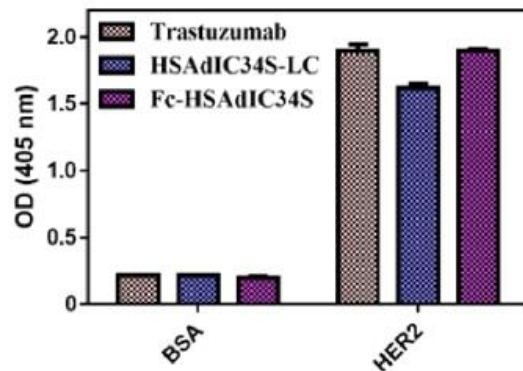
Chemical Lysine modification 2

- α , β -unsaturated sulfonamide type -

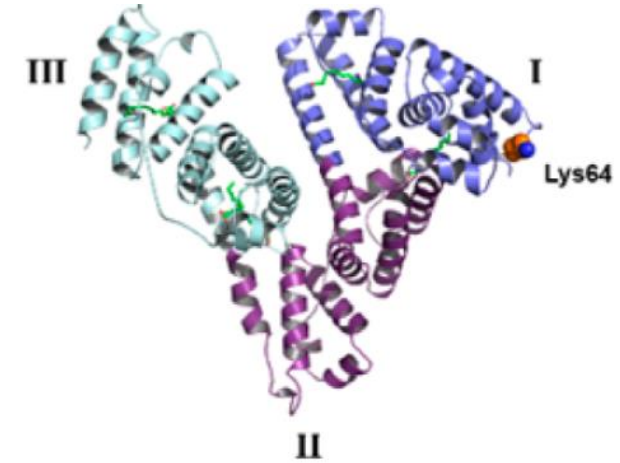
Time-course study of trastuzumab, HSAdIC34S-LC, and Fc-HSAdIC34S labeling with 10 equiv of cHx-Rho.



HER2 for trastuzumab and fusion conjugates.



Antibody conjugation using HSAdI as a fusion protein should be amenable to therapeutic applications.



Advantage

- ✓ Site-selectivity
- ✓ Antibody conjugates were also stable in human plasma

Disadvantage

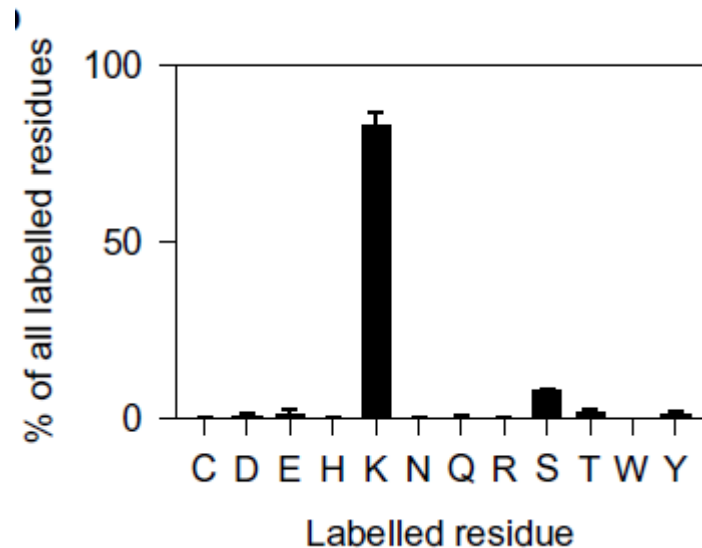
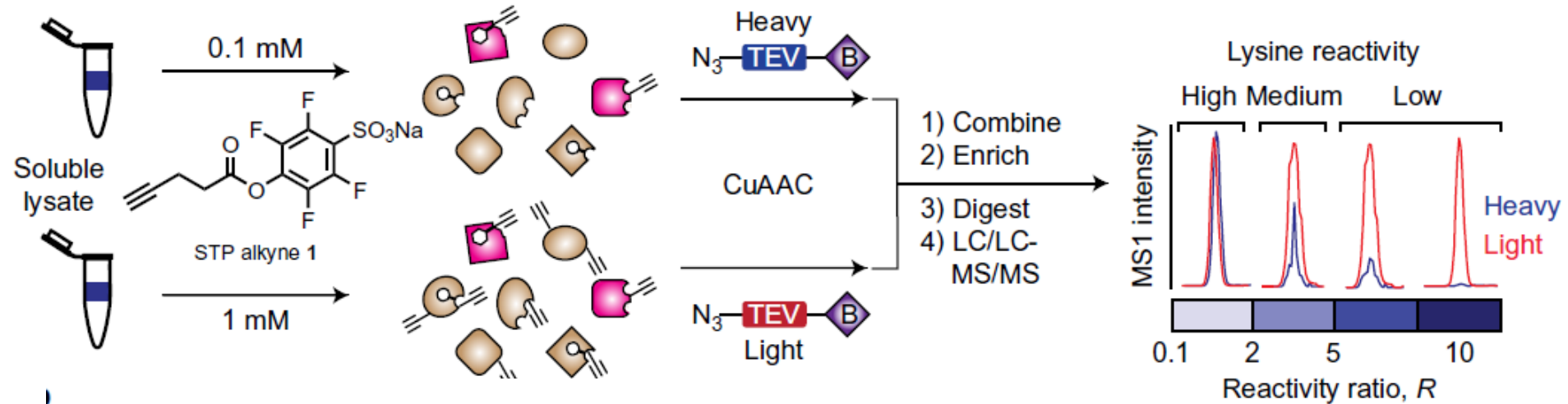
- ✓ Incomplete conversion
- ✓ Need large amount of reagent (10 eq)
- ✓ Basic condition. (pH=9.0)

Barbas et al., *Angew. Chem. Int. Ed.* **2014**, 53, 11783–11786

Barbas et al., *Bioconjugate Chem.* **2016**, 27, 2271–2275 16

Chemical Lysine modification 3

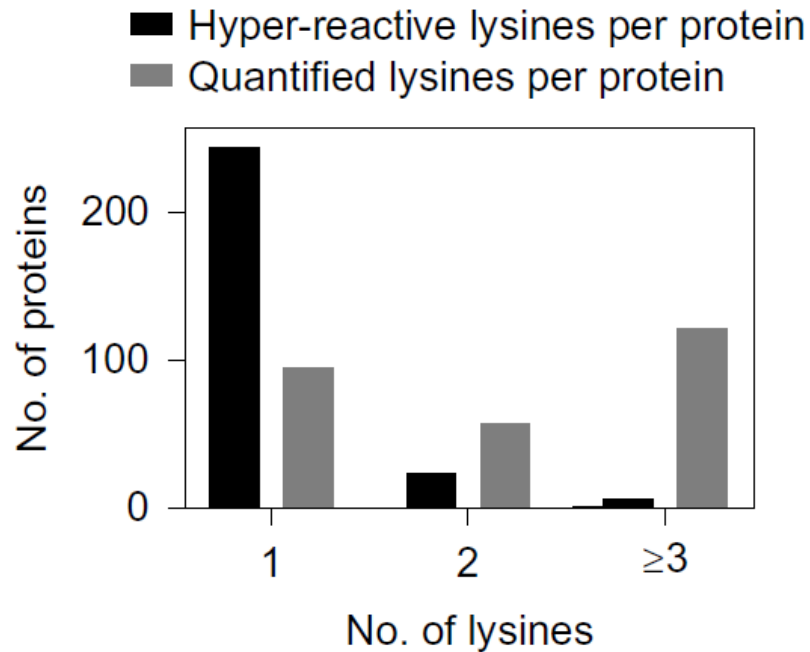
- sulfotetrafluorophenyl esters -



- ✓ broad reactivity
- ✓ good selectivity for lysine residues in the human proteome.

Chemical Lysine modification 3

- sulfotetrafluorophenyl esters -



Advantage

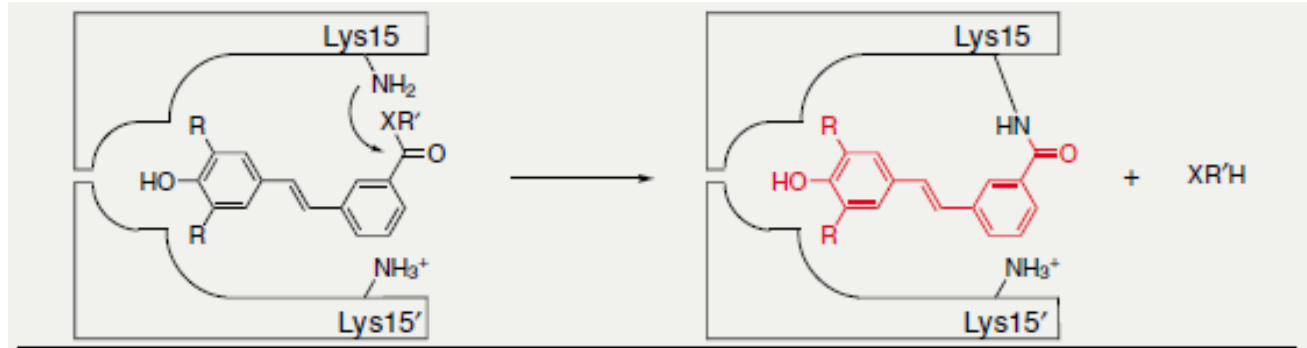
- ✓ more stable in aqueous solution than NHS esters.

Disadvantage

- ✓ the localization of hyper-reactive lysines to pockets could also restrict their access to post-translational machinery, such as ubiquitylation processes
- ✓ In order to prevent overreaction it is necessary to well control the equivalence of reagent and the reaction time.

Chemical Lysine modification 4

- Stilbene- *designed stilbenes that selectively and covalently modify the prominent plasma protein transthyretin*



Transthyretin(TTR) : One of the causative proteins of amyloidosis.

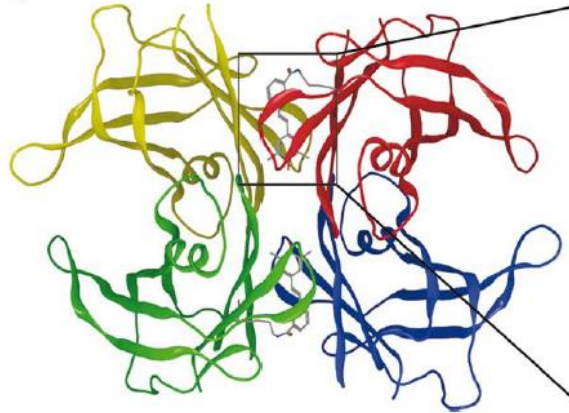


Figure. Crystal structure of the WT TTR

- **Amyloidosis caused by transthyretin** -

- Senile systemic amyloidosis (SSA)
- Familial amyloid polyneuropathy (FAP)

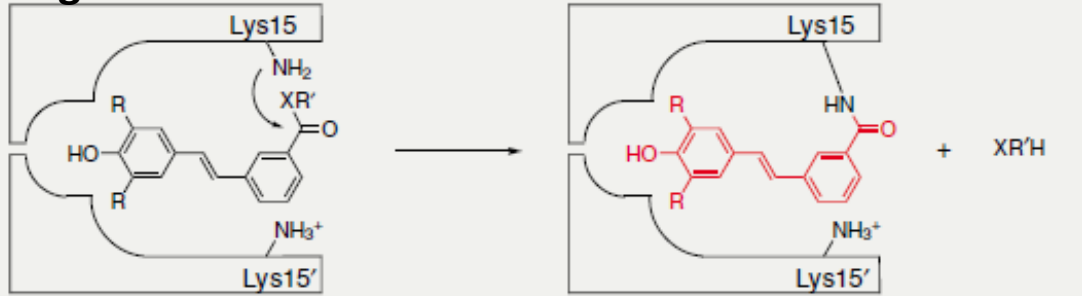
When transthyretin becomes unstable, it becomes amyloid fiber and aggregates.

➡ *A treatment method to stabilize transthyretin is required.*

Chemical Lysine modification 4

- Stilbene- designed stilbenes that selectively and covalently modify the prominent plasma protein transthyretin

Design of chemoselective covalent TTR kinetic stabilizers



Reduce electrostatic repulsion between lysine residues



Stabilization of transthyretin

Compound	R	XR'	% fibril formation ^a		IC ₅₀ (μM) ^b	% modification of TTR subunits in human blood plasma ^c
			7.2 μM	3.6 μM		
1	Br		2% (±1%)	12% (±1%)	2.00	36% (±0.4%)
			3% (±0.3%)	17% (±0.2%)		
2	CH ₃		2% (±1%)	16% (±2%)	2.00	48% (±1.3%)
			3% (±0.2%)	34% (±0.5%)		
3	CH ₃		4% (±1%)	21% (±2%)	2.26	49% (±0.7%)
			3% (±0.2%)	33% (±2%)		
4	CH ₃		1% (±1%)	16% (±1%)	1.96	32% (±3.3%)
			7% (±0.2%)	36% (±3%)		

Most chemoselective

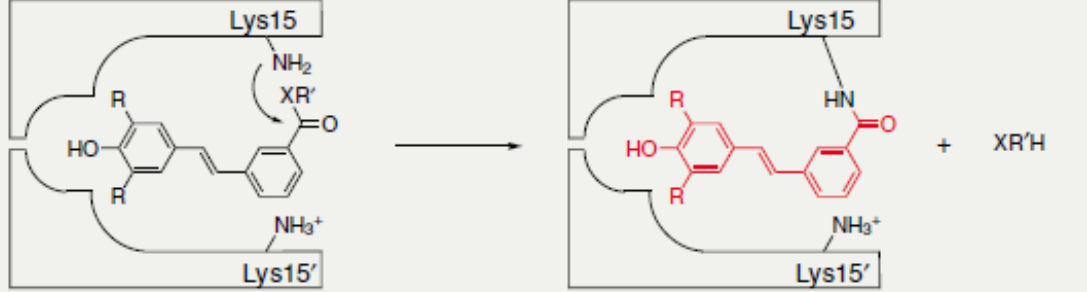
- Percent fibril formation of WT TTR(3.6 μM, black font) and V30M TTR(3.6 μM, blue font)
- Maximum modification percent is 50%.

- ✓ Ester or thioester group
→allow amino group of Lys to approach at Burgi-Dunitz angle

Chemical Lysine modification 4

- Stilbene- designed stilbenes that selectively and covalently modify the prominent plasma protein transthyretin

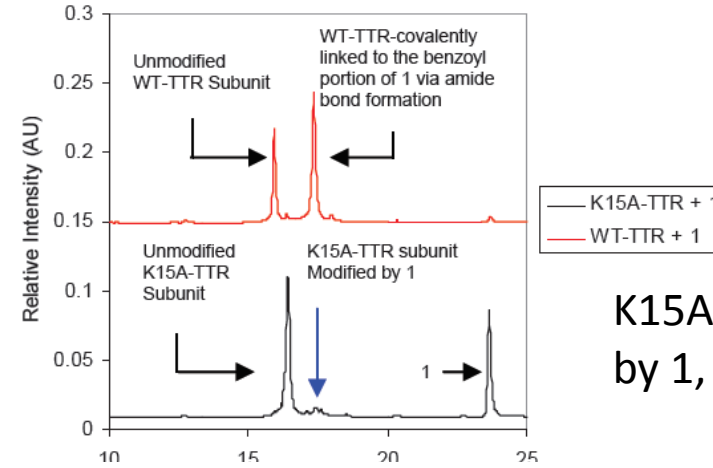
Design of chemoselective covalent TTR kinetic stabilizers



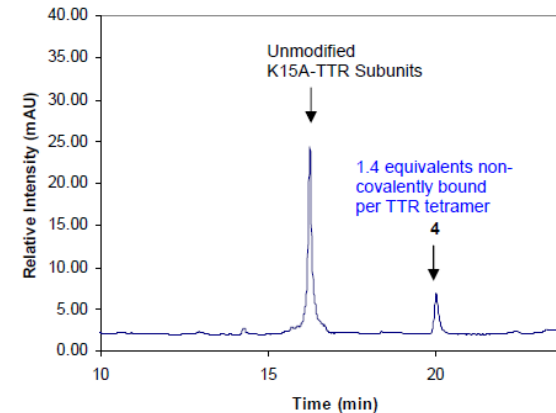
Compound	R	XR'	% fibril formation ^a		IC ₅₀ (μM) ^b	% modification of TTR subunits in human blood plasma ^c
			7.2 μM	3.6 μM		
1	Br		2% (±1%)	12% (±1%)	2.00	36% (±0.4%)
			3% (±0.3%)	17% (±0.2%)		
2	CH ₃		2% (±1%)	16% (±2%)	2.00	48% (±1.3%)
			3% (±0.2%)	34% (±0.5%)		
3	CH ₃		4% (±1%)	21% (±2%)	2.26	49% (±0.7%)
			3% (±0.2%)	33% (±2%)		
4	CH ₃		1% (±1%)	16% (±1%)	1.96	32% (±3.3%)
			7% (±0.2%)	36% (±3%)		

Most chemoselective

- Percent fibril formation of WT TTR(3.6 μM, black font) and V30M TTR(3.6 μM, blue font)
- Maximum modification percent is 50%.



K15A-TTR subunit modified by 1, 2, 3 was observed.

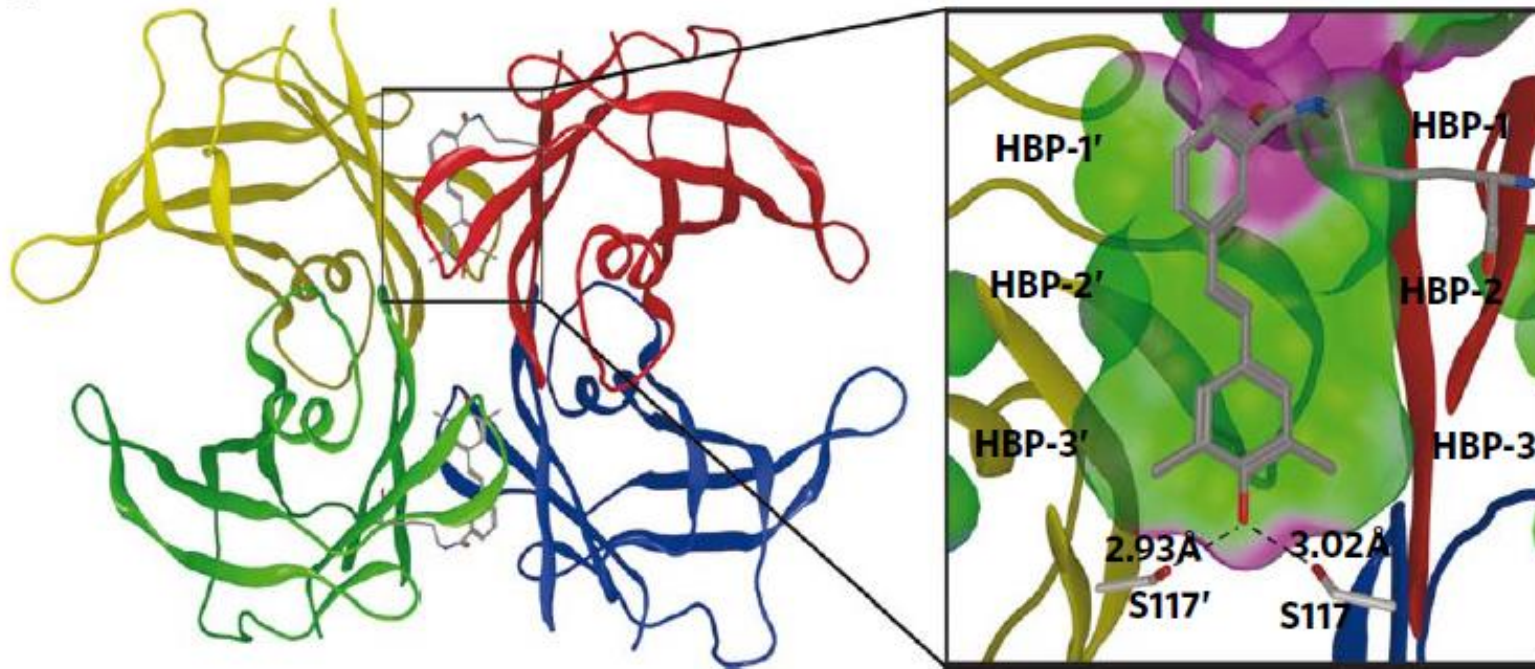


Not observed
→ High chemoselectivity

Chemical Lysine modification 4

- Stilbene- *designed stilbenes that selectively and covalently modify the prominent plasma protein transthyretin*

Transthyretin(TTR) : One of the causative proteins of amyloidosis.



Green: hydrophobic
Purple: polar
Blue: exposed

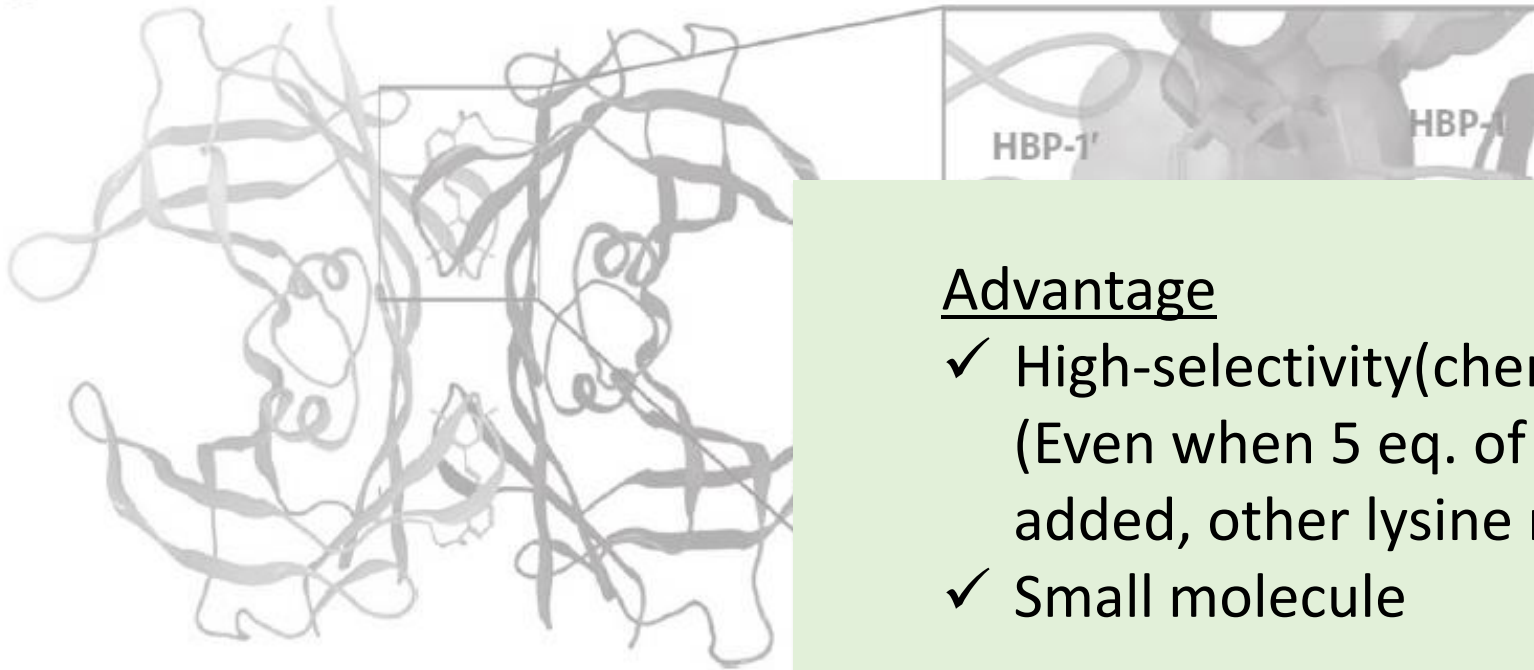
Figure. Crystal structure of the WT TTR

- ✓ Bridging hydrogen bonds are formed between the 4-OH of the benzoyl substructure and the Ser117 and Ser117' hydroxyls from adjacent TTR monomers.
- ✓ One Lys15 ϵ -amine group and one Lys15' ϵ -ammonium group at pH 7.

Chemical Lysine modification 4

- Stilbene- *designed stilbenes that selectively and covalently modify the prominent plasma protein transthyretin*

Transthyretin(TTR) : One of the causative proteins of amyloidosis.



Advantage

- ✓ High-selectivity(chemo-, site-)
(Even when 5 eq. of stilbene derivative were added, other lysine residues did not react)
- ✓ Small molecule

Disadvantage

- ✓ Applicable range is limited to TTR

- ✓ Bridging hydrogen bonds are formed between Ser117' hydroxyls from adjacent TTR subunits
- ✓ One Lys15 ϵ -amine group and one Lys117' ϵ -amine group are involved in the formation of the HBP-1

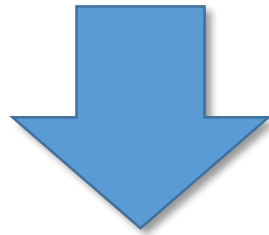
Chemical Lysine modification 5

- β -lactam- *The first site-specific ADC to be generated using a natural Lys for conjugation.*

ADC : useful for cancer treatment.

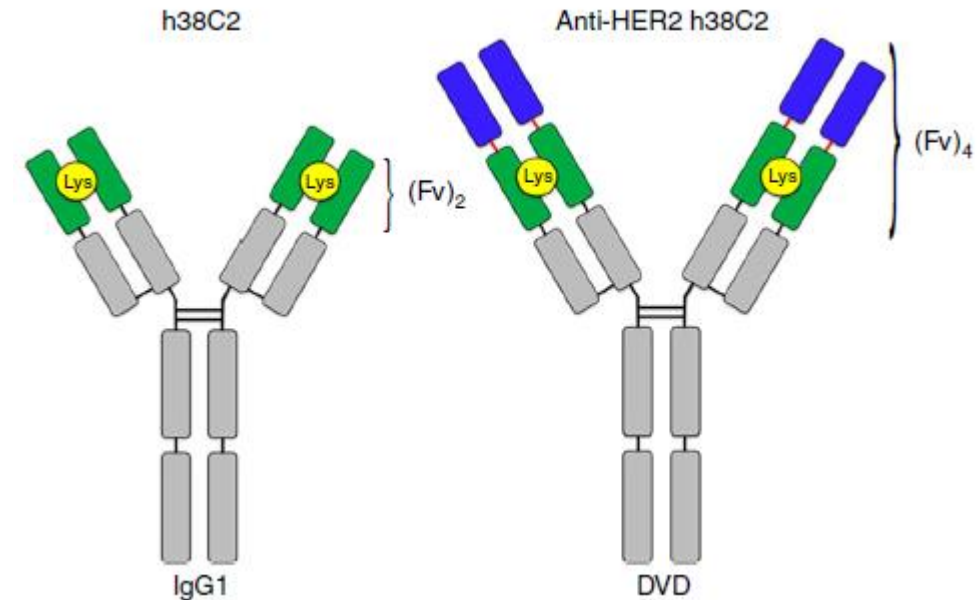
Conventional Site-Selective reaction

- ✓ unnatural amino acid by a genetic engineering technique
- ✓ introduce a sequence suitable for an enzymatic reaction



h38C2 :(humanized anti-hapten monoclonal antibody)

- ✓ Having a nucleophilic Lys residue ($pK_a \approx 6$) in the hydrophobic pocket



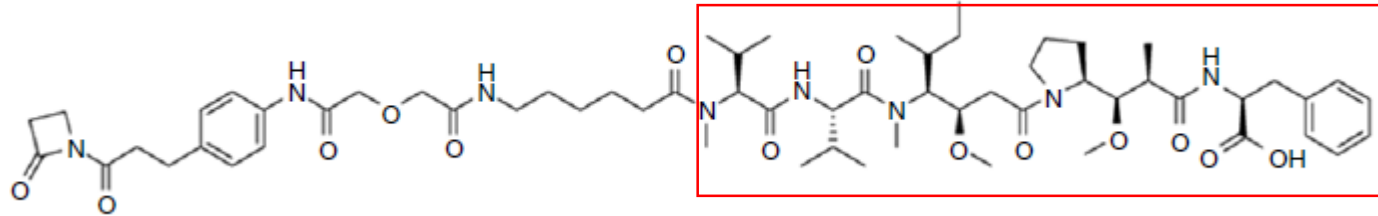
The DVD is composed of

- variable domains of trastuzumab (blue),
- h38C2 (green) with reactive Lys (yellow circle),
- constant domains (gray).

It becomes a selective modified target at in physiological pH.

Chemical Lysine modification 5

- β -lactam- *The first site-specific ADC to be generated using a natural Lys for conjugation.*



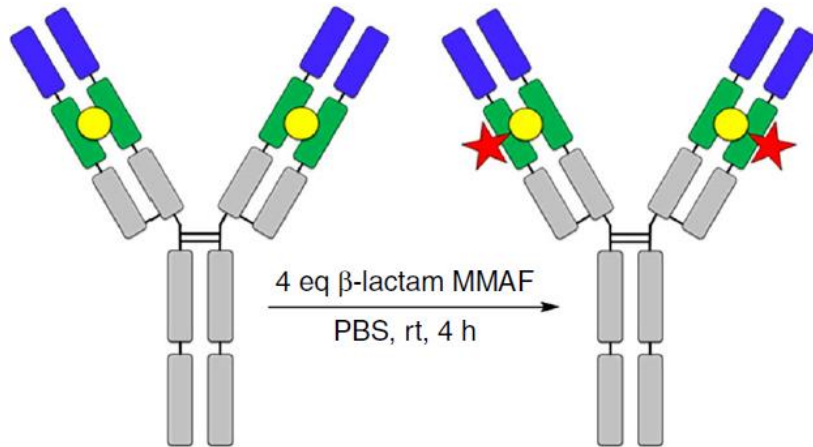
Monomethyl auristatin F (MMAF)
...To reduce cytotoxicity

Advantage

- ✓ Mutation free
- ✓ React irreversibly with Lys.
 - prevent premature drug release
- ✓ ADCs to be highly homogeneous
- ✓ conjugation does not eliminate any positive charges on the antibody
 - preserve electrostatic properties of the antibody

Disadvantage

- ✓ The possibility that other Lys react with reagent when the reaction time is longer.



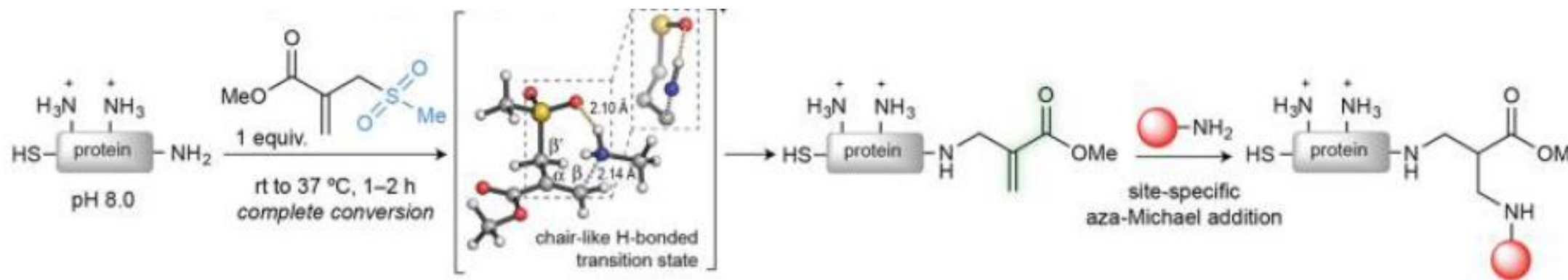
Contents

- Amino acid for bioconjugation
- **Chemical Lysine modification**
 - *N*-hydroxysuccinimide(NHS)-ester
 - α , β -unsaturated sulfonamide
 - Sulfotetrafluorophenyl esters
 - β -lactam conjugation
 - Sulfonyl acrylate (Lysine activated)
- Summary

Sulfonyl acrylate-mediated lysine modification

- Sulfonyl acrylate (Lysine activated) -

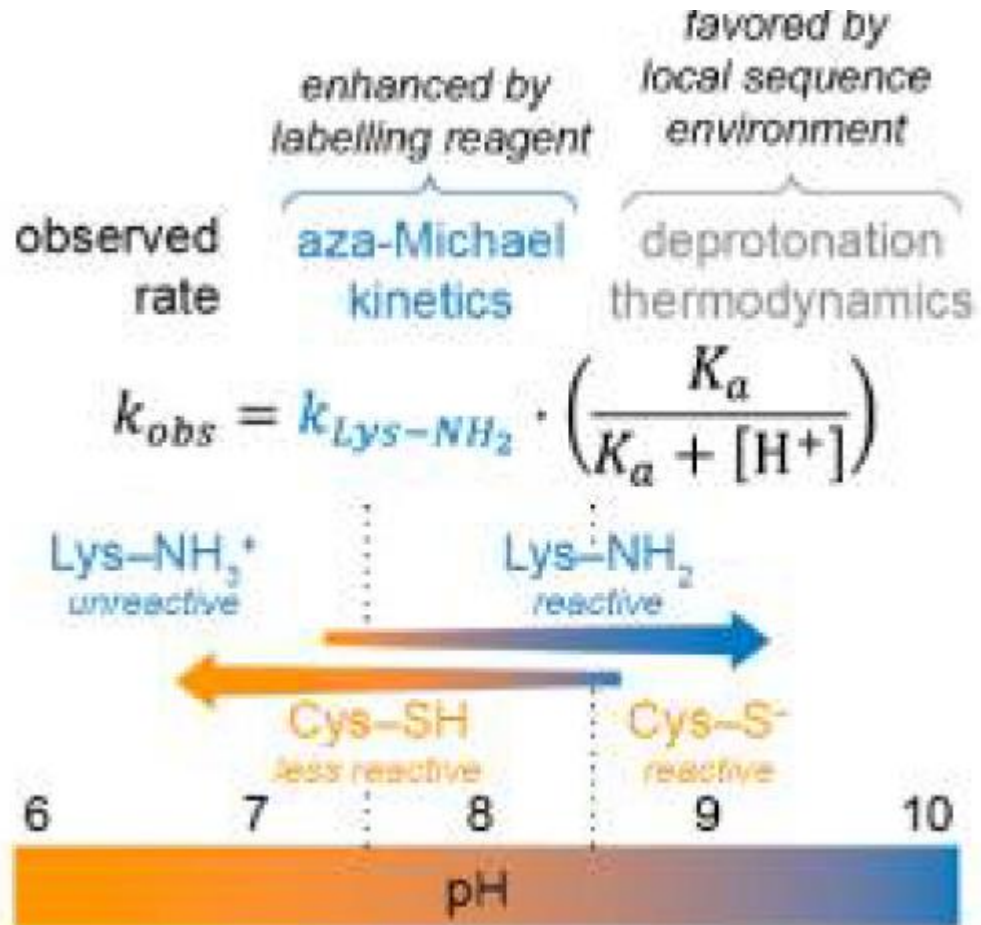
Hydrogen bond assisted chemo- and regioselective modification of lysine on native proteins



- ✓ Stoichiometric amount of sulfonyl acrylate reagent
- ✓ Proceed to completion rapidly (1~2h)
- ✓ Under mild conditions (pH=8.0, rt-37°C)
- ✓ Applicable to a range of native protein types
- ✓ The products are stable.
- ✓ The method is compatible with Cys bioconjugation.

pKa & reactivity of amino acids

- The reactivity of Lys and Cys residue at various pH conditions -



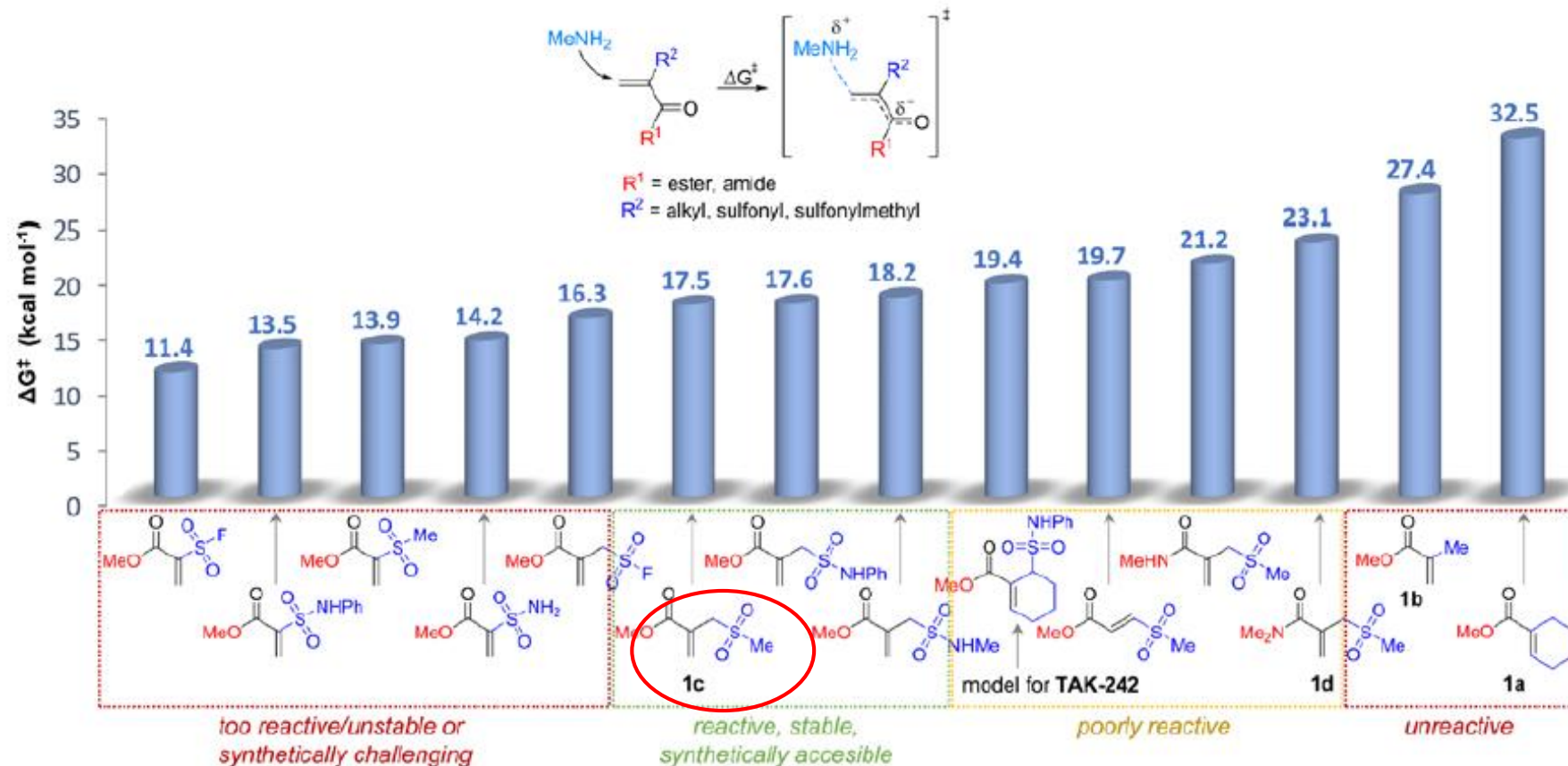
pH 5.5-6.0 Lysine residues are mostly protonated and unreactive.

pH 7.5-8.0 Lysine residues compete with cysteine as Michael donors.

pH 8.5- The more nucleophilic thiolate of cysteine residues usually dominate

Screening of Michael acceptors

- Computational screening of acrylic acid derivatives-



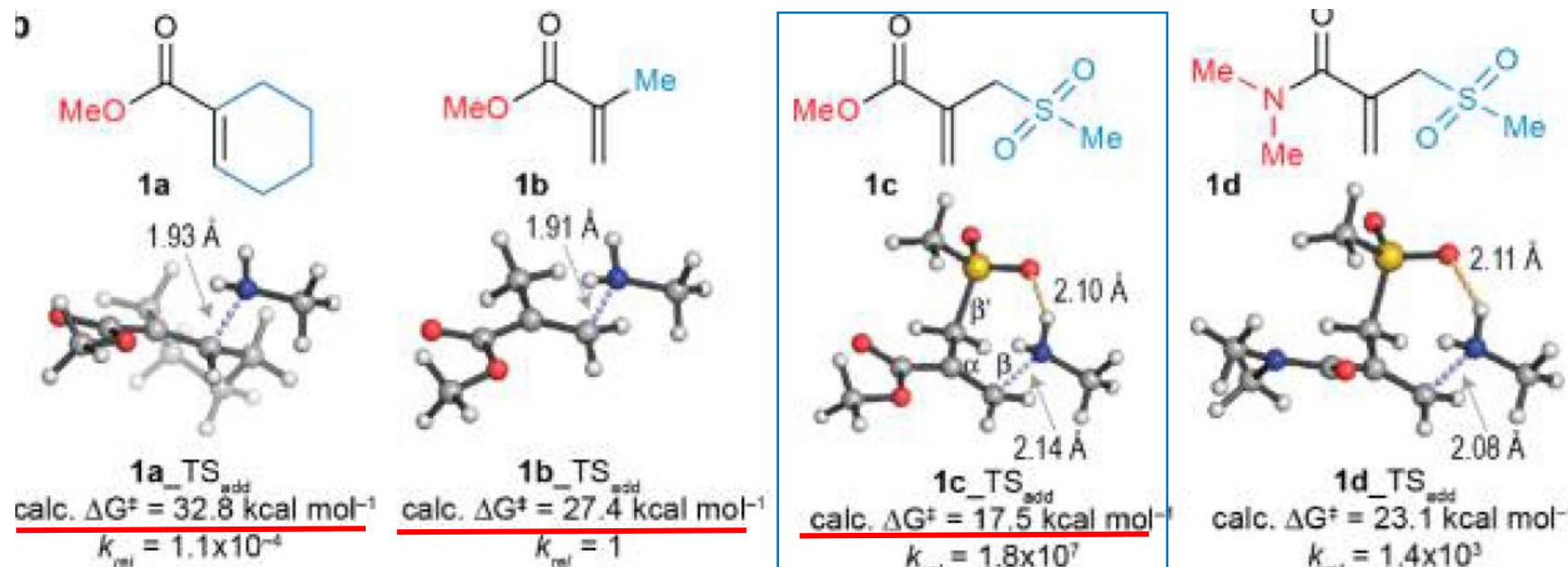
Activation barriers (ΔG^\ddagger) were calculated with PCM(H₂O)/M06-2X/6-31+G(d,p)

Sulfonylmethyl acrylate 1c was predicted to have a superior reactivity compared to its amide analogues

Detailed analysis of transition states

- Sulfonyl acrylate (Lysine activated) -

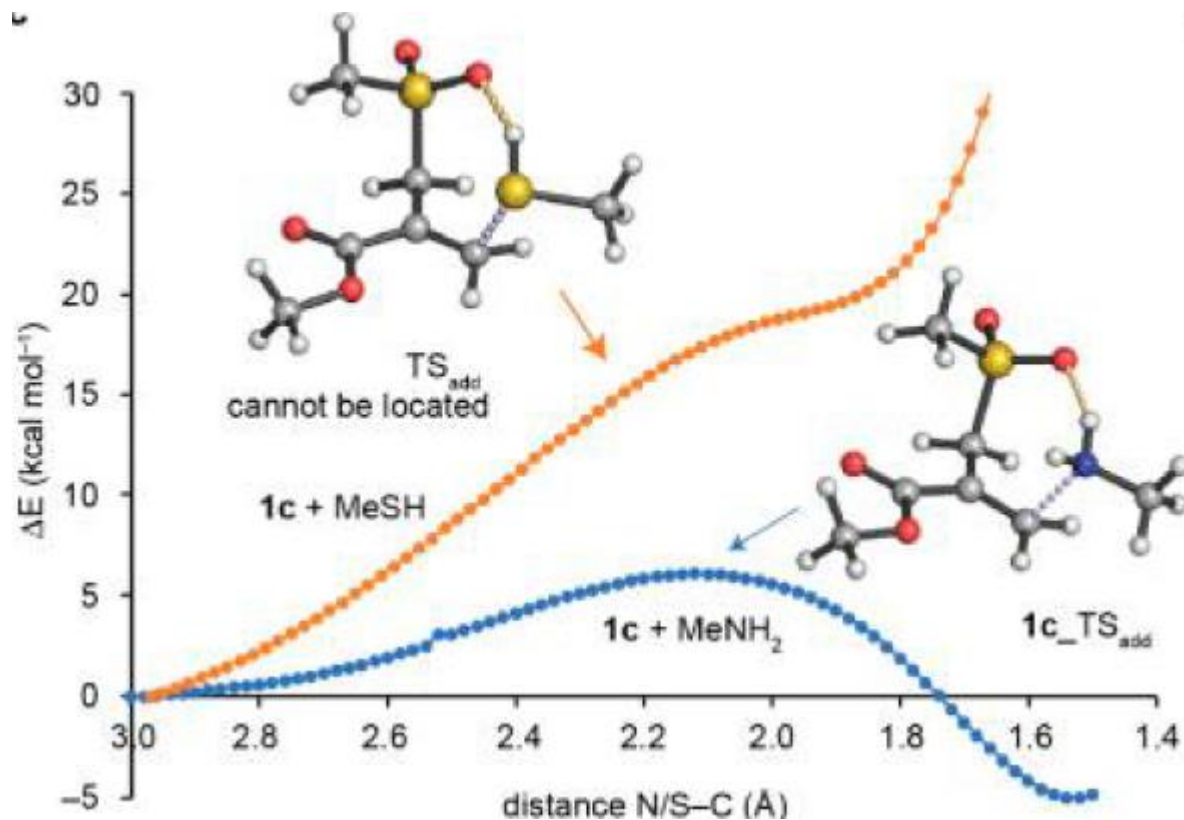
Acrylate electrophile derivatives 1a–d used in this study and transition states



This hydrogen bond interaction between the lysine model and the sulfone lowers the energy barrier by 10–16 kcal mol⁻¹

Comparison between Cys & Lys

- Sulfonylmethyl acrylate (Lysine activated) -



▪ Hydrogen-bonding

Cys < Lys

▪ Cys ... less polar character of the S-H bond,

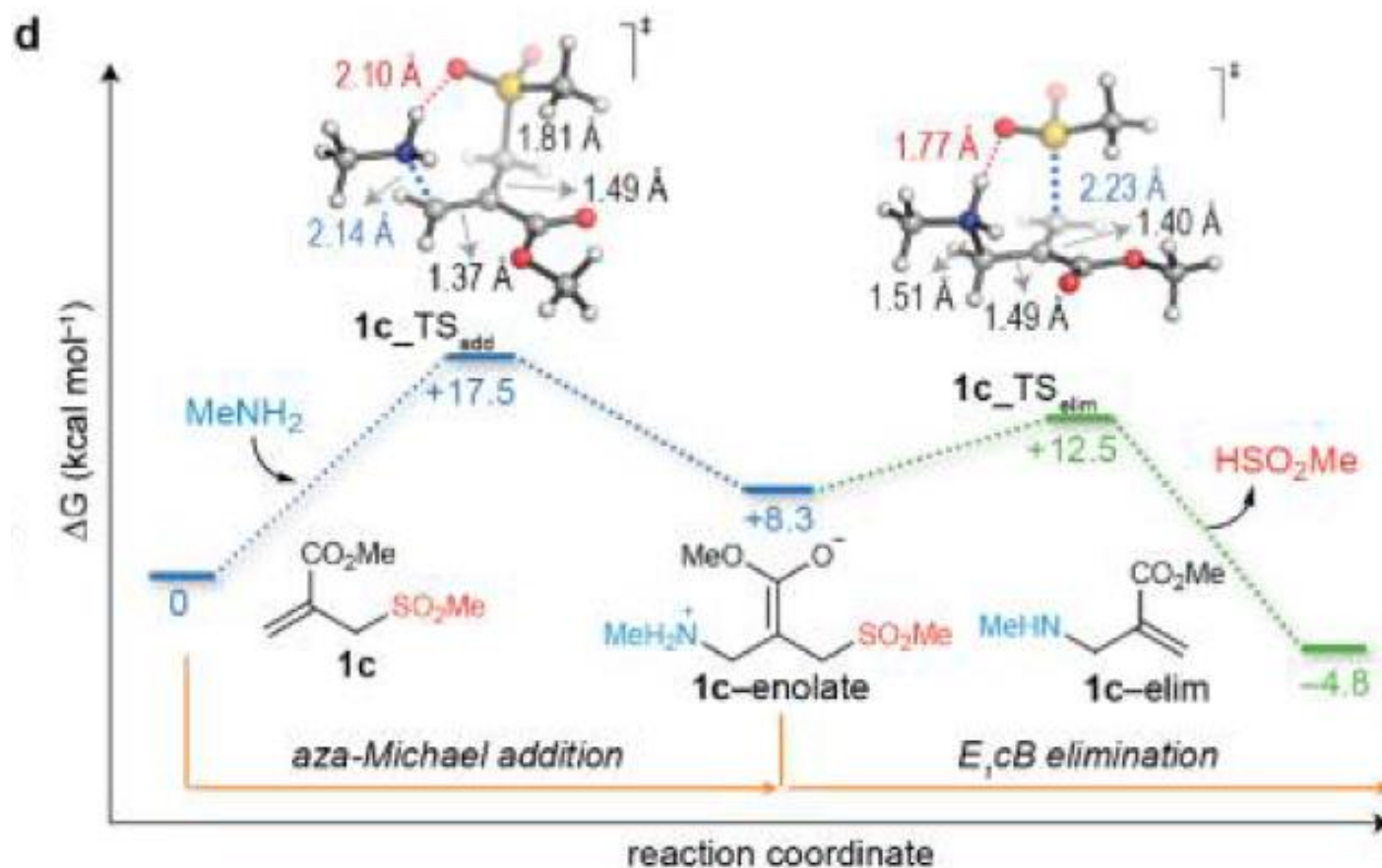


✓ *The positive charge developed at the amino group of the lysine model is efficiently dissipated by the sulfone*

A means to selectively modify lysine even in the presence of free cysteine residues at near neutral pH.

This conjugation is a two step reaction

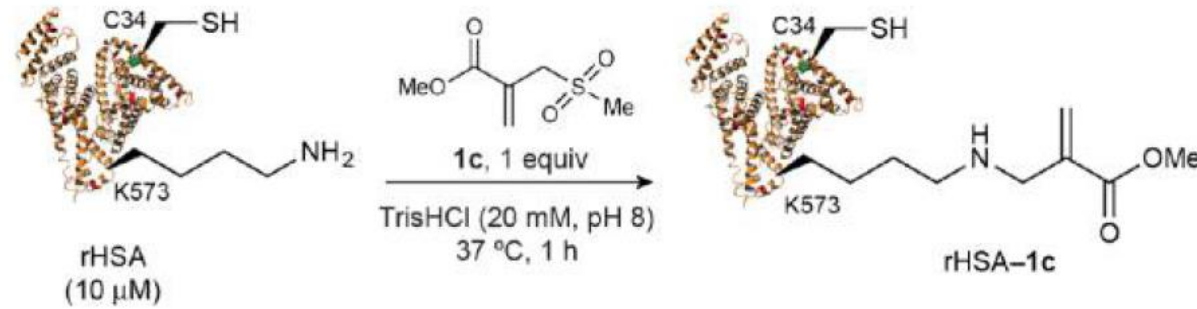
- Sulfonylmethyl acrylate (Lysine activated) -



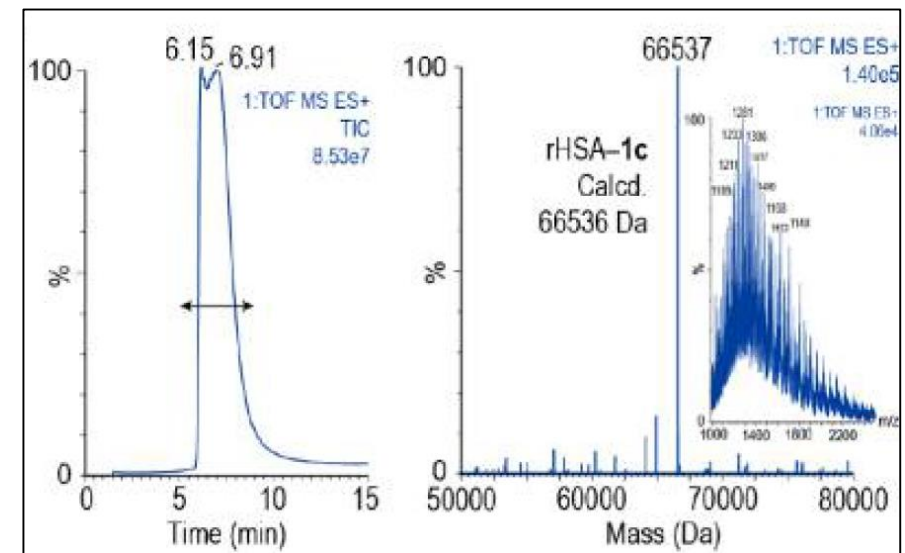
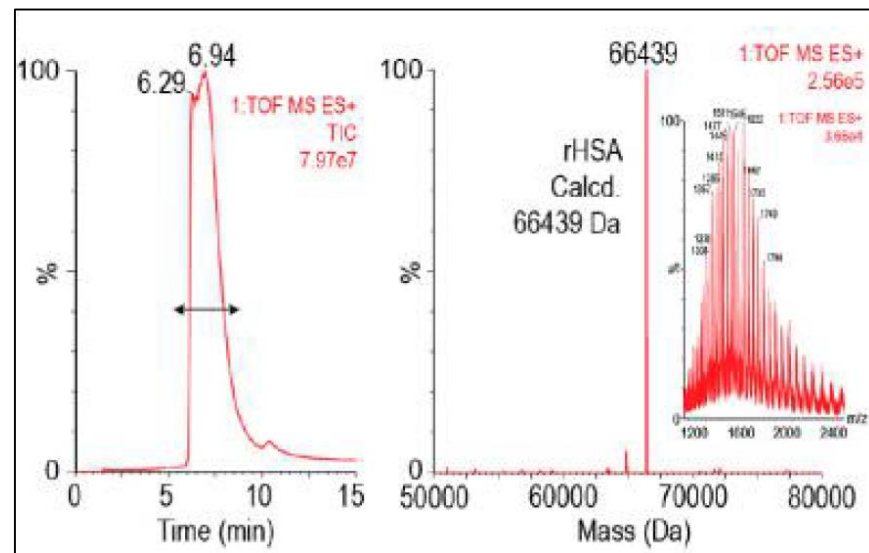
- ✓ Hydrogen bonding between the nucleophilic amino group and the sulfone moiety promotes both the aza-Michael addition and the elimination of methanesulfinic acid.

rHSA (MS analysis)

- Sulfonylmethyl acrylate (Lysine activated) -



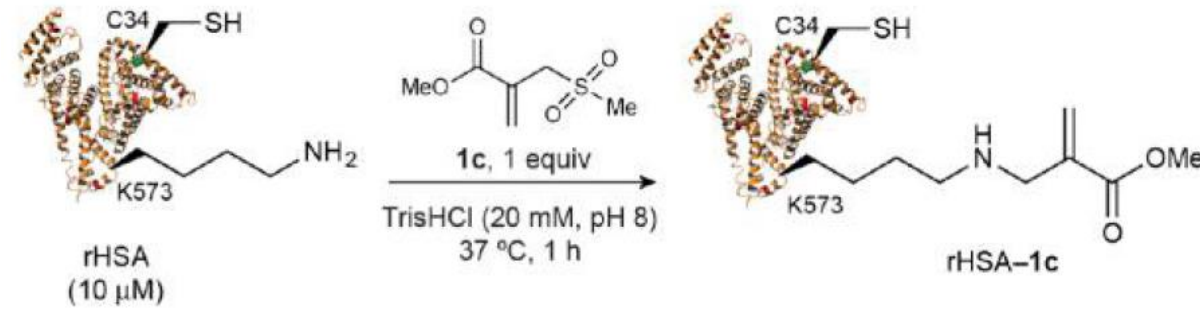
Total ion chromatogram



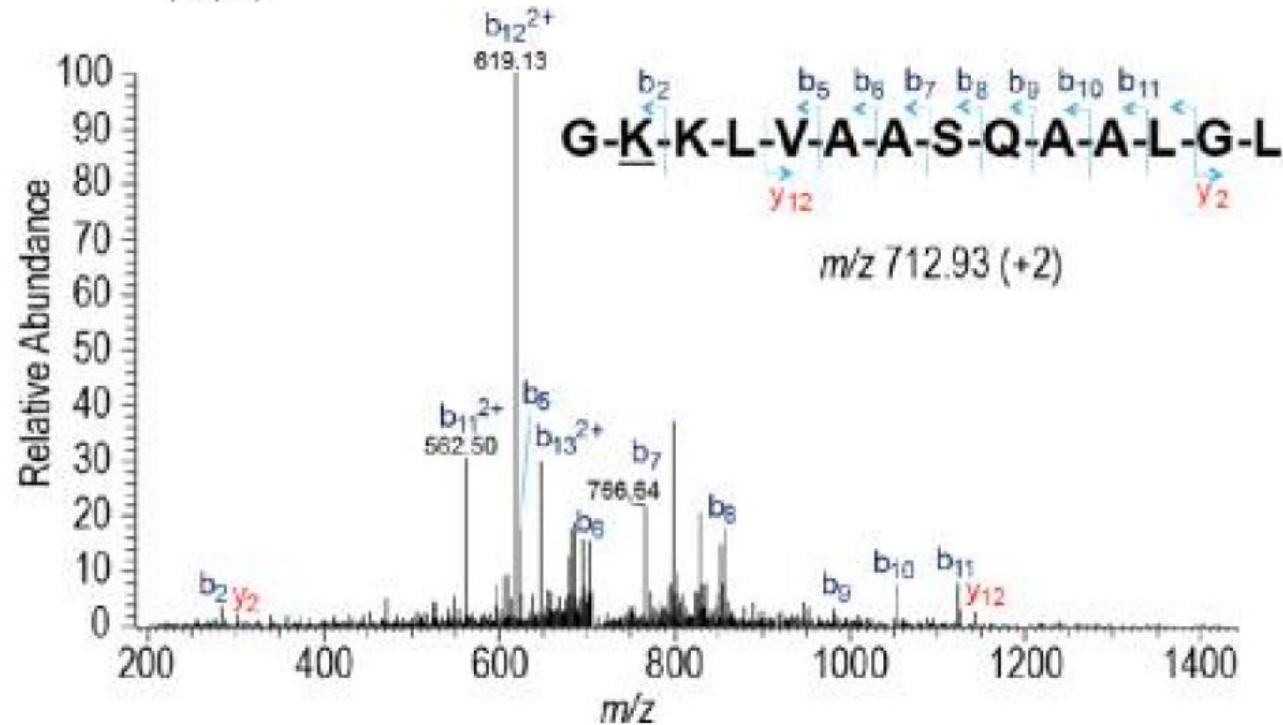
A single modification was produced in >95%

rHSA (LC-MS/MS, CD)

- Sulfonylmethyl acrylate (Lysine activated) -

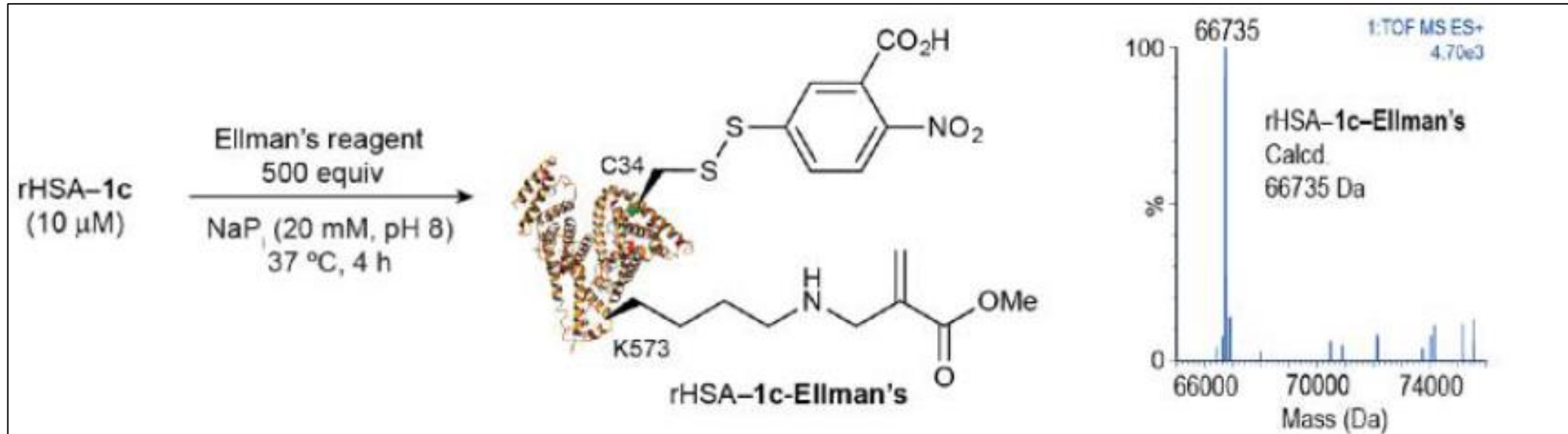


LC-MS/MS analysis

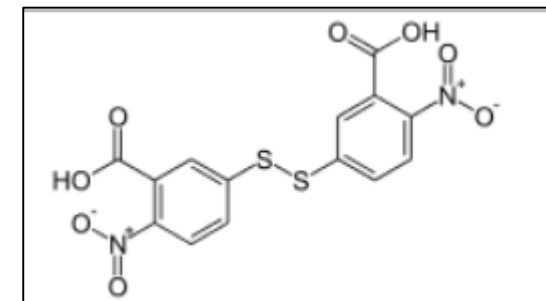


Reaction with thiol specific Ellman's reagent

- Sulfonylmethyl acrylate (Lysine activated) -



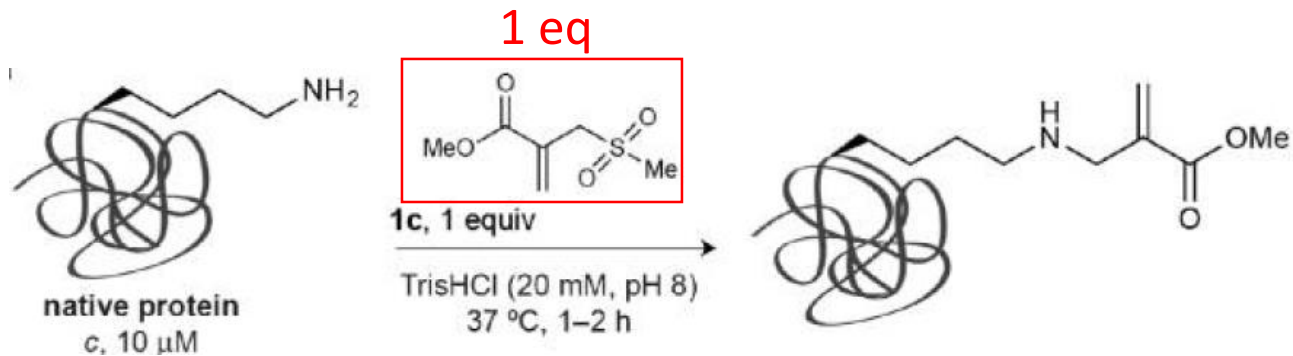
Ellman's reagent



- ✓ Fast
- ✓ Full conversion

Substrate scope

- Application to various type of native protein-



Conversion (%) native protein–1c (pH8, 37 °C) = 100

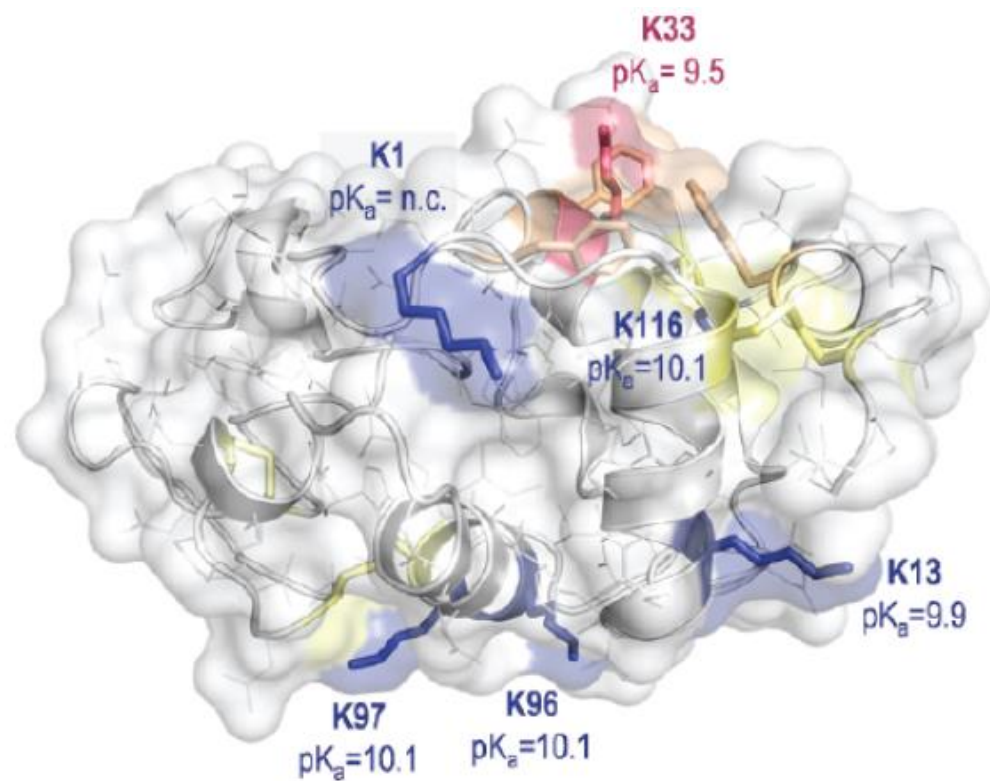
Conjugates are stable → Application to ADC is expected(Trastuzumab)

Regioselective lysine modification is applicable to a wide-range of native protein scaffolds.

Relationships between pKa & reactive site

- Sulfonyl acrylate (Lysine activated) -

Theoretical calculation of the most reactive lysine residue

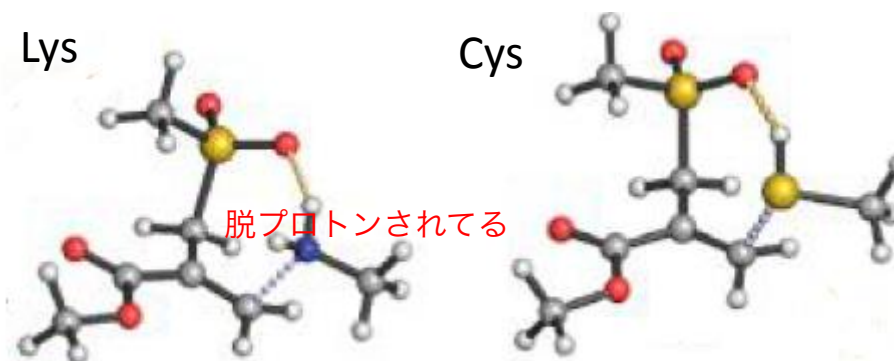


Lysozyme (6 Lys, no free Cys, 5 disulfides)

Predicted site of modification: K100
Observed site of modification: K100

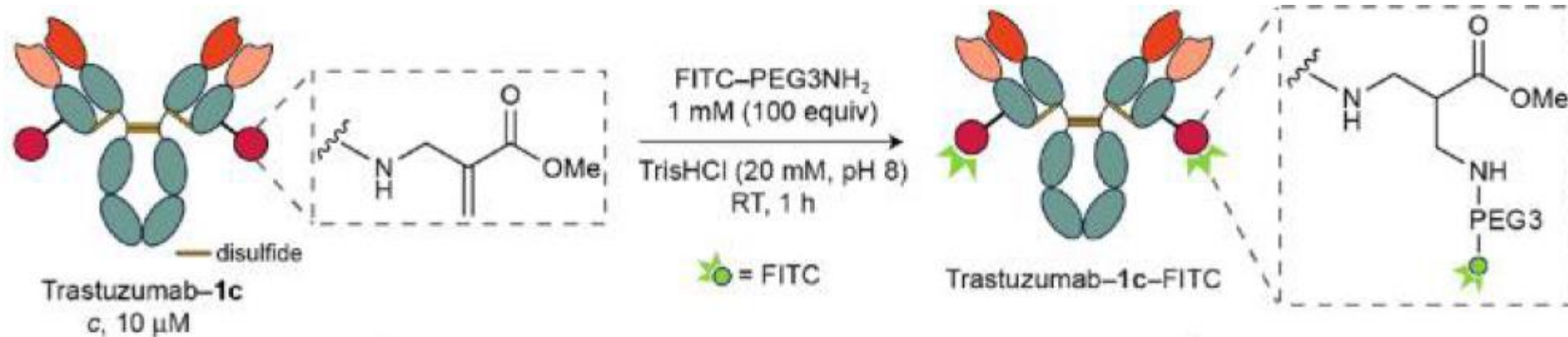
	pKa	modification
K100	10.1	observed
C95	10.3	Not observed

The selectivity of Lysine residue was observed because of Hydrogen bond.



Site-specific labeling

- The ability to precisely conjugate fluorophores and cytotoxic drugs to antibodies -



Conventional antibody labeling

... relied on labeling using disulfide bonds with **genetically engineered** free lysine residues or **non-natural amino acids**.

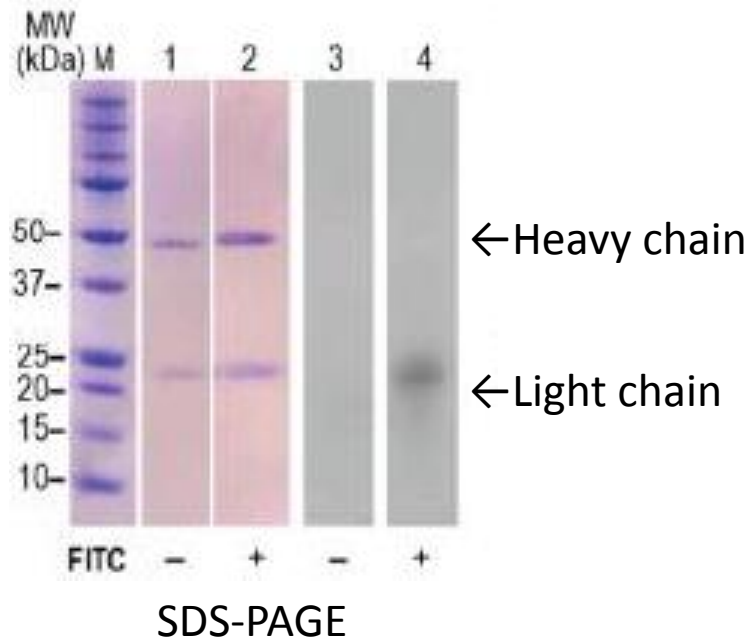
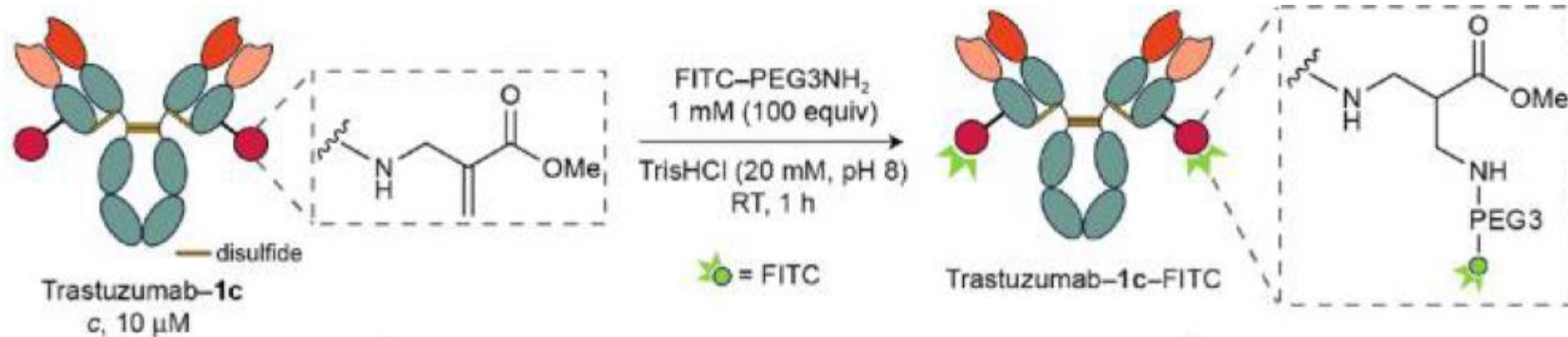
→ Produce heterogeneous compounds

There are possibility that affinity with the target antigen may be compromised.

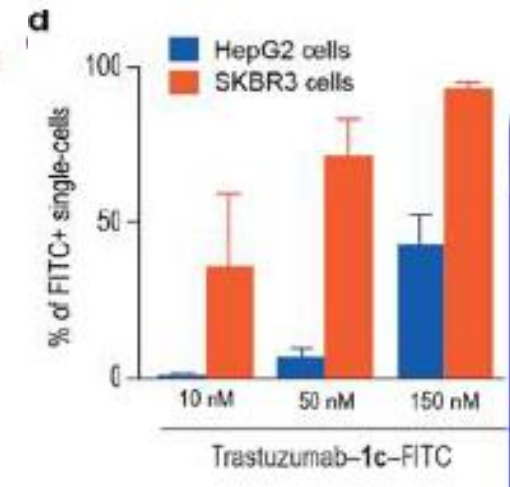
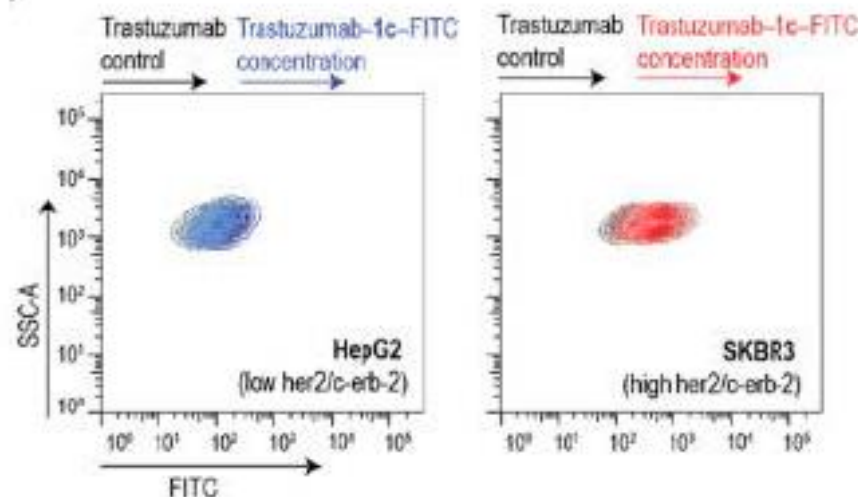
This method can be applied directly to a therapeutic antibody in its native form.

Site-specific labeling

- The ability to precisely conjugate fluorophores and cytotoxic drugs to antibodies -



Flow cytometry



Contents

- Amino acid target for bioconjugation
- Chemical Lysine modification
 - *N*-hydroxysuccinimide (NHS)-ester
 - α , β -unsaturated sulfonamide
 - Sulfotetrafluorophenyl esters
 - β -lactam
 - Sulfonyl acrylate (Lysine activated)
- **Summary**

Summary

- Lysine residues are abundant and easy to target because of their high reactivity.
- A general way to selectively target single Lysine is lacking.
- This method can provide a single lysine one step modification with complete chemo- and region-selectivity.
- Direct applicability to wild-type protein sequence bode well for routinely accessing site-selectively modified proteins for basic biology and therapeutic applications.

Appendix

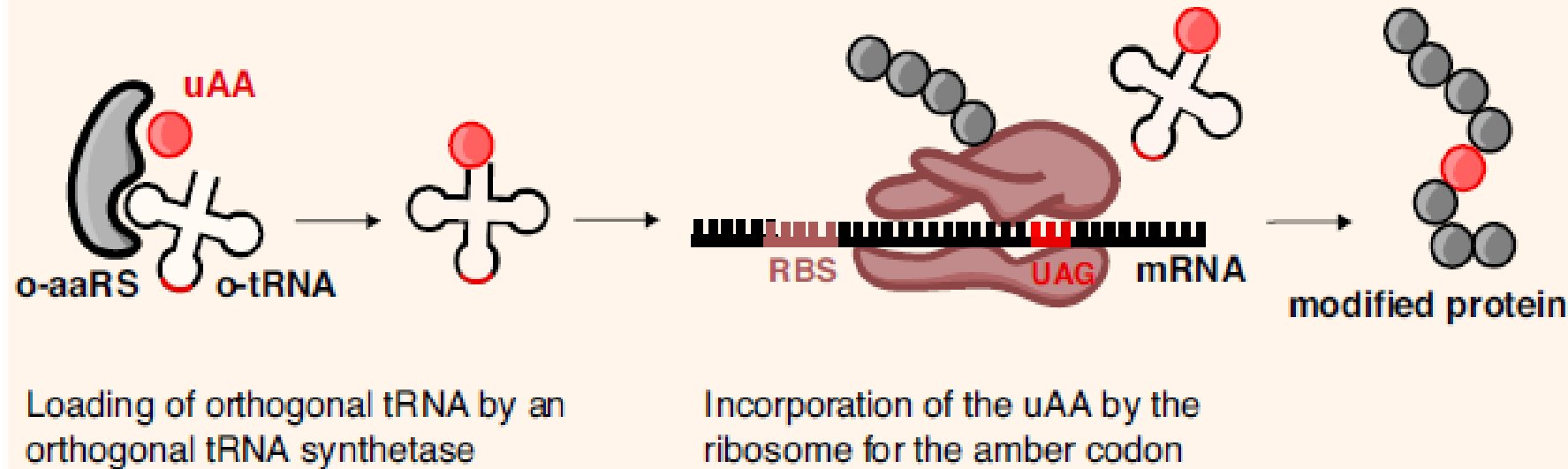
Various types of modification

- Ligation
- Genetic Code expansion
- Ligand – directed type
- Chemical conjugation using Dha

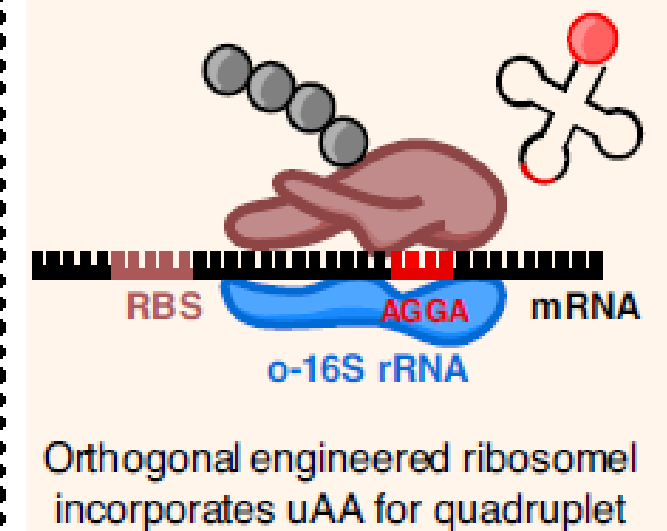
Various types of modification

- Genetic code expansion

Amber suppression

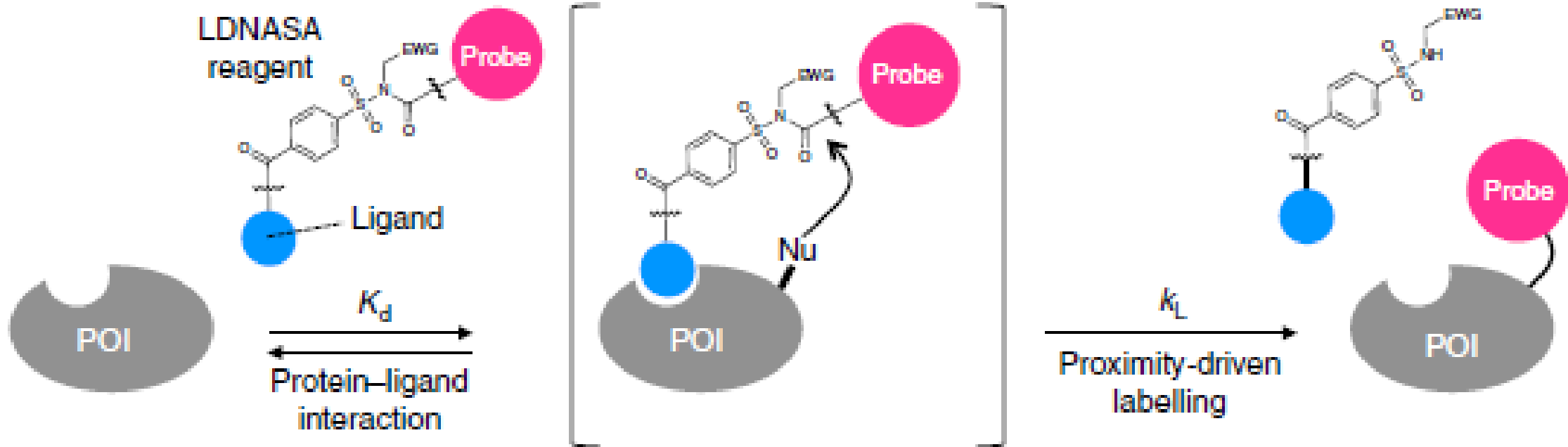


Quadruplet suppression



Various types of modification

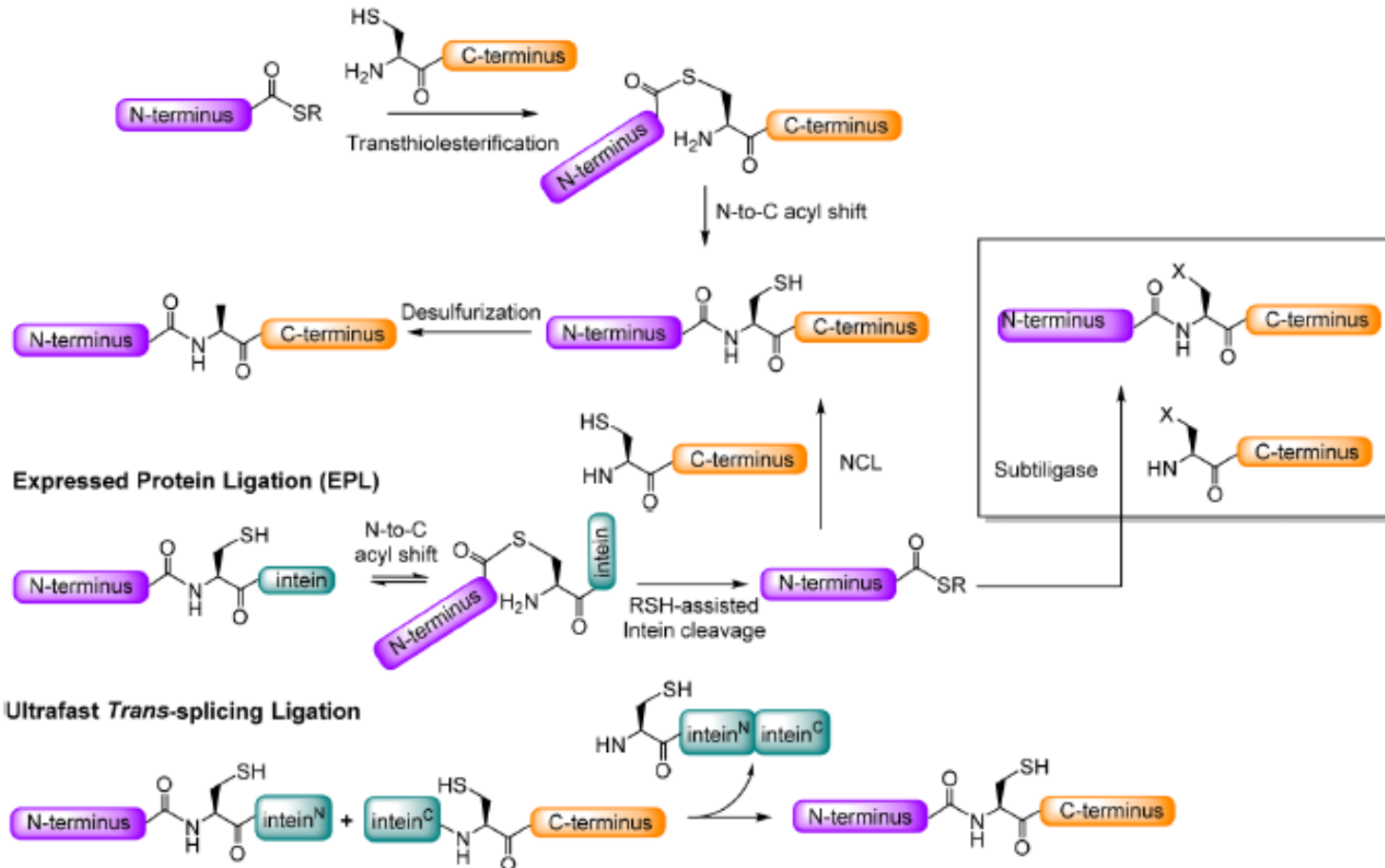
- Ligand-directed type



Various types of modification

- Ligation

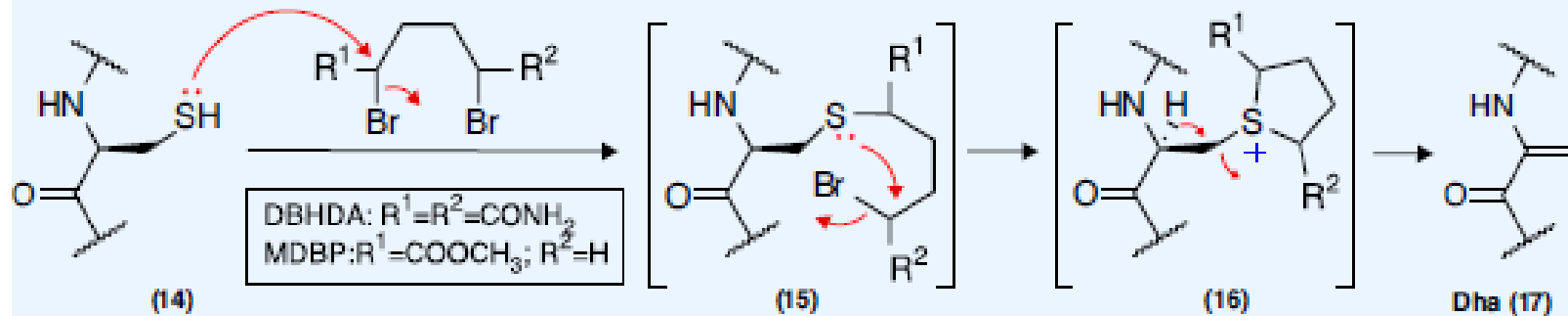
Native Chemical Ligation (NCL)



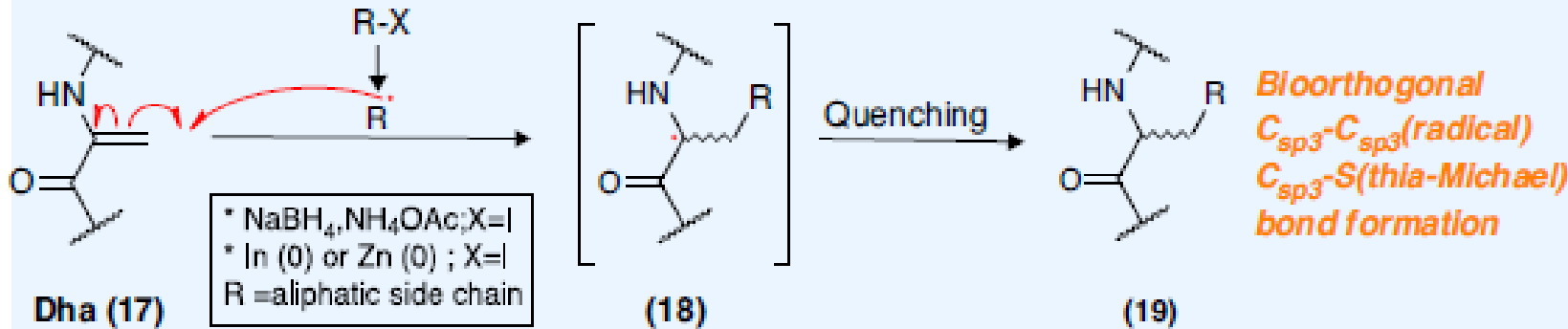
Various types of modification

- Chemical conjugation using Dha

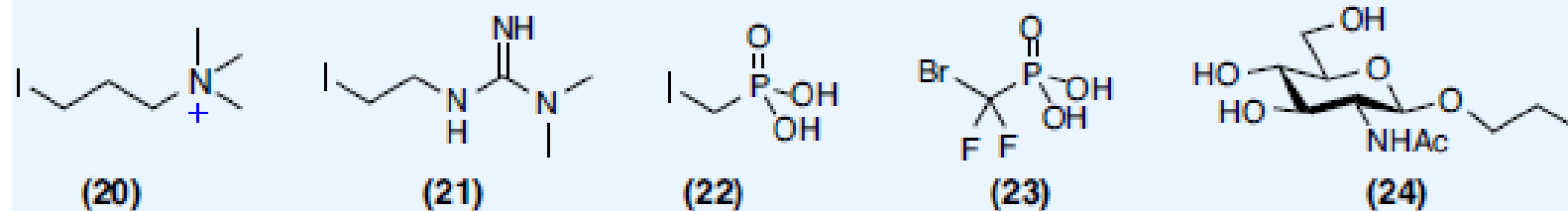
1. Bis-alkylation-elimination of cysteine to dehydroalanine (Dha)



2. Addition to dehydroalanine

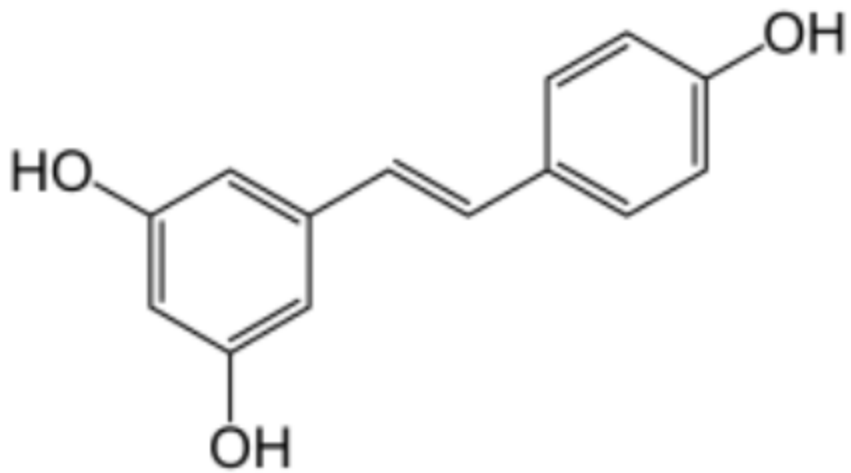


Examples of reagents used for chemical mutagenesis on histones:



Chemical Lysine modification 4

- Stilbene- *designed stilbenes that selectively and covalently modify the prominent plasma protein transthyretin*

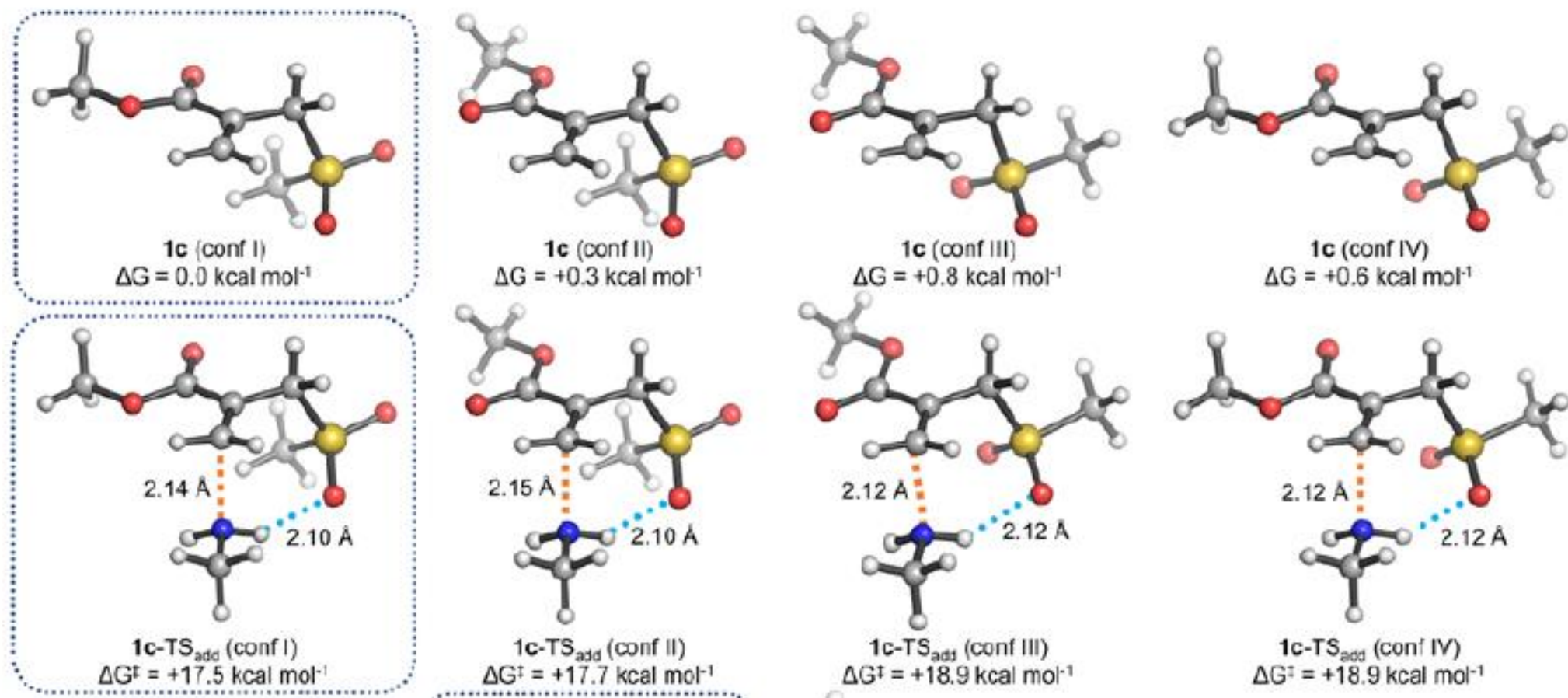


Resveratrol

- ✓ Noncovalent TTR kinetic stabilizers
- ✓ It is known to prevent amyloid formation– associated cytotoxicity, whereas structurally related compounds with poor TTR binding capacity do not inhibit cytotoxicity

Chemical Lysine modification 6

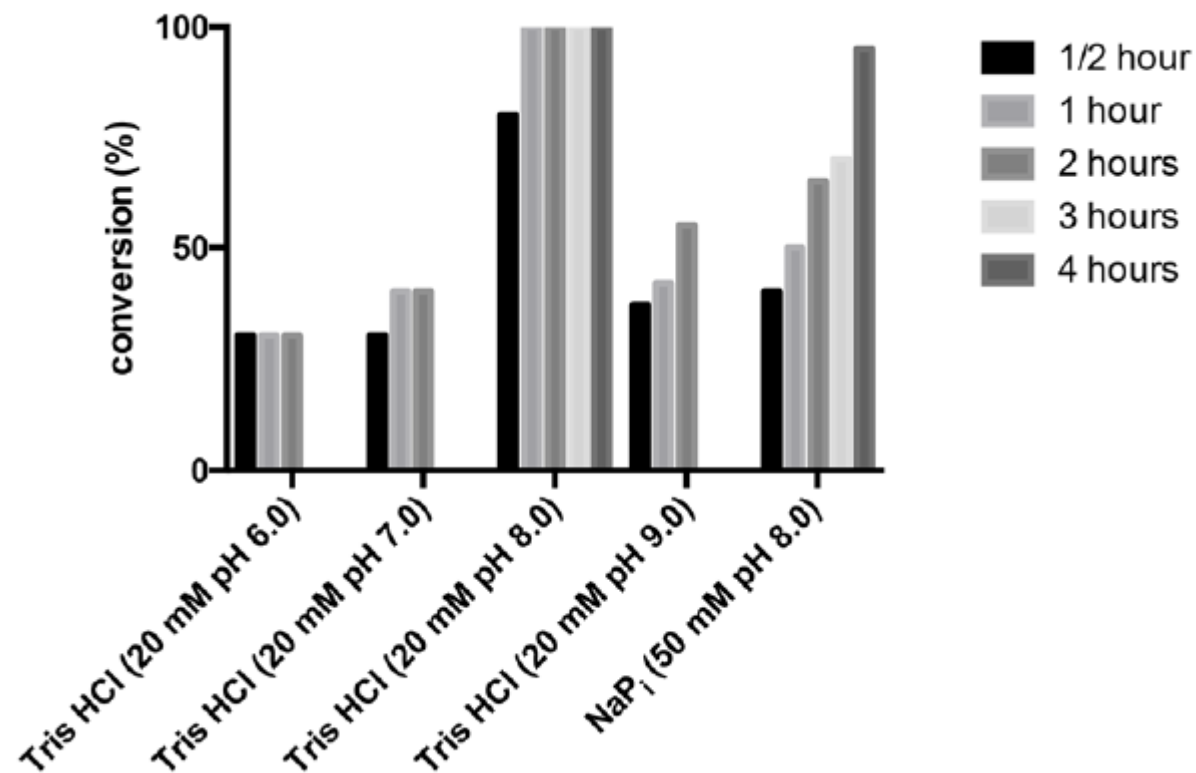
- Sulfonyl acrylate (Lysine activated) -



Chemical Lysine modification 6

Optimisation of reaction conditions between rHSA and **1c** with respect to pH, buffer and time

Reaction conditions		Conversion (%) rHSA-1c (10 μ M, 37 $^{\circ}$ C)				
Buffer	pH	Time (h)				
		1/2	1	2	3	4
Tris HCl 20 mM	6.0	30	30	30	-	-
	7.0	30	40	40	-	-
	8.0	80	100	100	100	100
	9.0	37	42	55	-	-
NaPi 50 mM	8.0	40	50	65	70	95



Chemical Lysine modification 3

Target protein: **Lysozyme**

Production run: **40 ns**

Protonation state change attempted every **5** simulation steps

pH	5.0	6.0	7.0	8.0	9.0	10.0	11.0	12.0	13.0	14.0
LYS13	0.999993	0.999947	0.997636	0.992302	0.945720	0.398367	0.188996	0.012162	0.001551	0.000046
LYS33	0.999820	0.999562	0.997475	0.949310	0.775744	0.208294	0.032031	0.004438	0.000303	0.000018
LYS96	0.999989	0.999846	0.998788	0.990135	0.900597	0.547084	0.099162	0.011172	0.001762	0.000174
LYS97	0.999971	0.999822	0.997938	0.978243	0.900403	0.554339	0.091912	0.011426	0.001499	0.000092
LYS116	0.999996	0.999961	0.998962	0.986439	0.898668	0.542344	0.079672	0.010680	0.000642	0.000143

Protonated fraction ($1 - f_d$) for each residue as a function of the pH