

## Allyl Sulfides Are Privileged Substrates in Aqueous Cross-Metathesis: Application to Site-Selective Protein Modification

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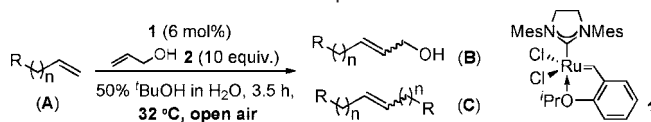
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Olefin metathesis has become a mainstay in organic synthesis.<sup>1</sup> Cross-metathesis (CM), however, is largely underdeveloped compared to ring closing metathesis (RCM) and ring opening metathesis polymerization since CM does not have the entropic driving force of RCM and is complicated by self-metathesis.<sup>2</sup> Our group has a long-term interest in site-selective chemical modification of proteins in an effort to study and modulate their function.<sup>3</sup> Olefin metathesis is an attractive way to install these protein modifications through a stable carbon–carbon bond. Indeed, incorporation of olefins into proteins has been possible for nearly a decade,<sup>4</sup> but metathesis at such residues has not been realized. Despite recent reports of olefin metathesis in water, the current benchmark for homogeneous aqueous CM is the self-metathesis of simple unsaturated alcohols such as allyl alcohol.<sup>5,6</sup> The limited examples revealed to date highlight the challenges for aqueous CM and the gap in substrate complexity that must be bridged to carry out metathesis on protein surfaces.

To determine the viability of CM on protein surfaces, simple amino acid models were investigated. Substrates were selected on the basis of potential incorporation into proteins. A reasonable starting point was homoallylglycine (Hag) since its *in vivo* incorporation by methionine auxotrophic *Escherichia coli* is known.<sup>4</sup> Hoveyda–Grubbs second generation catalyst **1**<sup>7</sup> was selected since it is phosphine free and therefore more likely to be compatible with protein disulfides than other conventional catalysts. A simple test metathesis with allyl alcohol **2** was carried out to assess the reactivity of Hag derivative **3**. At the outset, we limited ourselves to temperatures generally compatible with proteins ( $\leq 37$  °C) and made no effort to exclude oxygen. Since **1** is not freely soluble in water, it was added as a solution in *t*-BuOH. Unfortunately, despite repeated attempts, only starting material **3** was recovered (Table 1, Entry 1). We turned next to cysteine derivatives since incorporation into proteins should be possible by either chemical or genetic means if they proved reactive in CM. Remarkably, *S*-allylcysteine (Sac) derivative **4** underwent metathesis with allyl alcohol (Entry 2), affording the CM product in 56% isolated yield (74% based on recovered **4**). This result was noteworthy given the number of instances where thioethers were detrimental to ruthenium-based metathesis catalysts.<sup>8</sup> The metathesis was also efficient with allyl homocysteine **5** and bisamide Sac derivative **6**. Yet when the alkene was extended by one or two methylene units from the sulfur center, only allyl alcohol self-metathesis was observed along with recovered starting material (Entries 5 and 6). Other allylheteroatom substrates were screened, but allyl sulfides remained the most efficient metathesis substrates under the conditions employed (Entries 7–13).

While the self-metathesis of allyl sulfides has been carried out in organic solvents, the efficiency relative to other heteroatom or hydrocarbon analogues was not apparent and yields were highly dependent on the catalyst used.<sup>9</sup> We suggest that the enhanced reactivity of allyl sulfides may be a consequence of sulfur

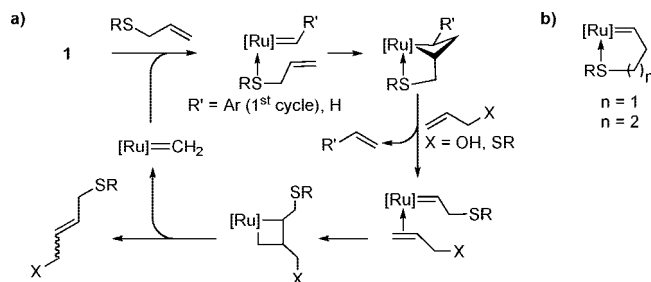
**Table 1.** Heteroatom Effects in Aqueous Cross-Metathesis



Entry	Alkene <b>A</b>	Cross-Metathesis <b>B</b> (%) <sup>c</sup>	Self-Metathesis <b>C</b> (%) <sup>c</sup>
1 <sup>a,b</sup>		3	0
2 <sup>a</sup>		4	56 (74 brsm)
3 <sup>a</sup>		5	67 (99 brsm)
4 <sup>a</sup>		6	68 <sup>d</sup>
5 <sup>a,b</sup>		7	0
6 <sup>a</sup>		8	0
7		9	0
8		10	52 <sup>d</sup>
9		11	19
10		12	8
11		13	11
12		14	31
13		15	28 <sup>d</sup>

<sup>a</sup> 8 mol % of **1**, 2.5 h. <sup>b</sup> 30% *t*-BuOH/H<sub>2</sub>O. <sup>c</sup> Isolated. <sup>d</sup> >98% conversion.

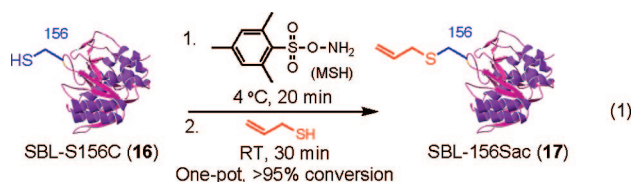
### Scheme 1. Sulfur Assisted Cross-Metathesis of Allyl Sulfides



coordination to the ruthenium center that brings the reacting centers into close proximity (Scheme 1a).<sup>10</sup> Fürstner and co-workers have noted a similar “relay effect” of appropriately positioned heteroatoms in RCM macrocycle synthesis.<sup>11</sup> Vinyl sulfides are poor substrates in aqueous metathesis, likely leading to Fischer carbenes sensitive to water (Table 1, Entry 7). The decreasing reactivity of

butenyl and pentenyl sulfides (e.g., **7–8** and **11–12**) may be attributed to the formation of unproductive five- or six-membered chelates, respectively (Scheme 1b).<sup>8d,12</sup> Attempts to isolate such species, however, were unfruitful.<sup>13</sup>

The results in Table 1 led us immediately to pursue Sac incorporation into proteins. Conveniently, an efficient chemical route to thioether modified proteins was recently developed in our laboratory.<sup>14</sup> The reaction of *O*-mesitylenesulfonylhydroxylamine (MSH) with cysteine rapidly generates dehydroalanine which can then be reacted with a thiol nucleophile. Application of this methodology to a single cysteine mutant of the serine protease subtilisin *Bacillus lentus* (SBL) allowed efficient incorporation of Sac into the protein (eq 1).



Ready access to Sac on protein surfaces enabled us to take advantage of the unique reactivity of allyl sulfides in CM. Initial attempts were carried out simply by adding excess **1** and **2** to a solution of SBL-156Sac **17** in 50 mM sodium phosphate (pH 8.0) (Table 2). LC-MS analysis revealed largely unreacted **17**, even after prolonged reaction time. Nevertheless, we were intrigued by a minor, yet significant, MS signal that appeared upon the addition of **1** to **17**.<sup>15</sup> We speculated this species might be a metalloprotein derived from metathesis with **1**, inactive in CM due to nonproductive chelation of side chains to ruthenium. MgCl<sub>2</sub> was added to the reaction buffer with the intention of disrupting any such nonproductive chelation to ruthenium. Fürstner used Ti(O<sup>i</sup>Pr)<sub>4</sub> in a similar fashion to disrupt nonproductive chelation in RCM.<sup>12c</sup> Gratifyingly, when MgCl<sub>2</sub> was included in the buffer, CM with allyl alcohol proceeded to >90% conversion at room temperature (Table 2, Entry 2).<sup>15</sup> To verify that the effect was due to Mg<sup>2+</sup> and not chloride, NaCl was used as additive (Table 2, Entry 3): no CM was observed

**Table 2.** Cross-Metathesis on SBL-156Sac

Entry	Alkene (mM)	Additives (mM)	Temp.	Prod.	Conversion (%) <sup>a</sup>
1	HO-CH <sub>2</sub> -CH=CH <sub>2</sub> ( <b>2</b> ) (100)	None	RT	-	0
2	<b>2</b> (100)	MgCl <sub>2</sub> (100)	RT	<b>18</b>	>90
3	<b>2</b> (100)	NaCl (100)	RT	-	0
4 <sup>b</sup>	<b>23</b> (30)	MgCl <sub>2</sub> (80)	37 °C	<b>19</b>	50
5 <sup>b</sup>	<b>24</b> (130)	MgCl <sub>2</sub> (130)	37 °C	<b>20</b>	60
6 <sup>c</sup>	<b>25</b> (75)	MgCl <sub>2</sub> (160)	37 °C	<b>21</b>	55
7 <sup>c</sup>	<b>26</b> (66)	MgCl <sub>2</sub> (130)	37 °C	<b>22</b>	60

<sup>a</sup> Determined by LC-MS. <sup>b</sup> First hour at rt. <sup>c</sup> First 2 h at RT.

without Mg<sup>2+</sup>. Importantly, **18** was an active peptidase and not denatured over the modification sequence.<sup>15</sup> Biologically and therapeutically relevant glycosylation<sup>16</sup> and PEGylation<sup>17</sup> were also achieved by CM (Entries 4–7).

Finally, efforts in genetic incorporation of allyl sulfide containing amino acids are also underway to explore their scope as tags for CM on proteins. Genetic installation ensures stereochemical homogeneity of the protein backbone and allows strategic flexibility. This approach was tested using the B834 *E. coli* strain, a methionine (Met) auxotroph.<sup>4,15</sup> Low level Sac incorporation was verified by MS–MS analysis in a single Met mutant of *Sulfolobus solfataricus* β-glycosidase expressed in Met-depleted media with Sac as Met surrogate.<sup>15</sup>

In conclusion, we have shown that allyl sulfides are effective substrates in aqueous CM through the use of catalyst **1**. Taking advantage of the enhanced reactivity of allyl sulfides in CM, we were able to post-translationally modify proteins via carbon–carbon bond formation. This work is an addition to a growing interest in metal-mediated protein modifications<sup>18</sup> and a new standard in substrate sensitivity and complexity in olefin metathesis.

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**Supporting Information Available:** Full experimental details and compound characterization. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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