

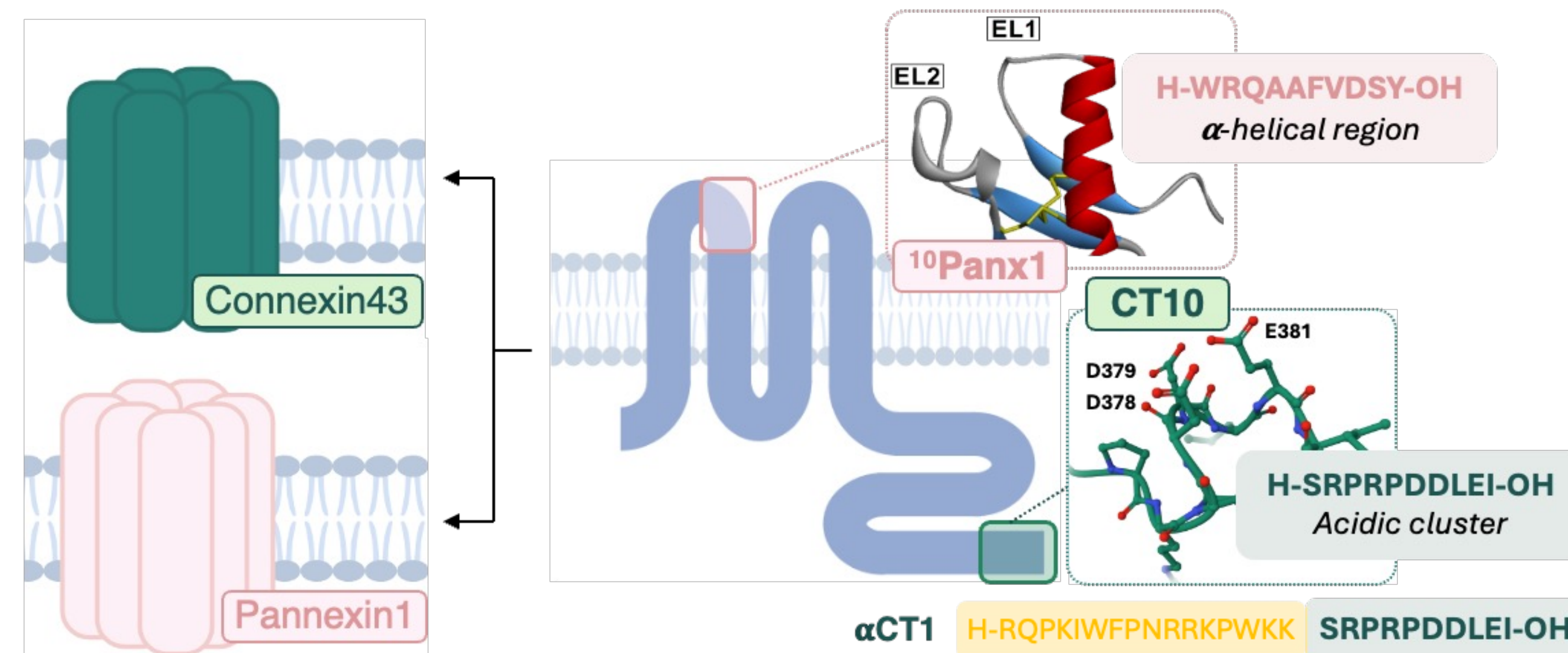
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CONNEXIN43 AND PANNEXIN1

Connexins (Cx) and Pannexins (Panx) are ubiquitous channel-forming membrane proteins which play an important role in a variety of both physiological and pathological processes. In the cardiac system, Connexin43 (Cx43) hemichannels (HCs) and Pannexin1 (Panx1) channels have shown to be critical mediators of inflammation in the context of **cardiovascular diseases**, including **ischemia-reperfusion (I/R) injury** and **atherosclerosis**. Selective modulation of Cx43 HC and Panx1 channels represents then a potential therapeutic approach.¹



CT10 AND ¹⁰Panx1

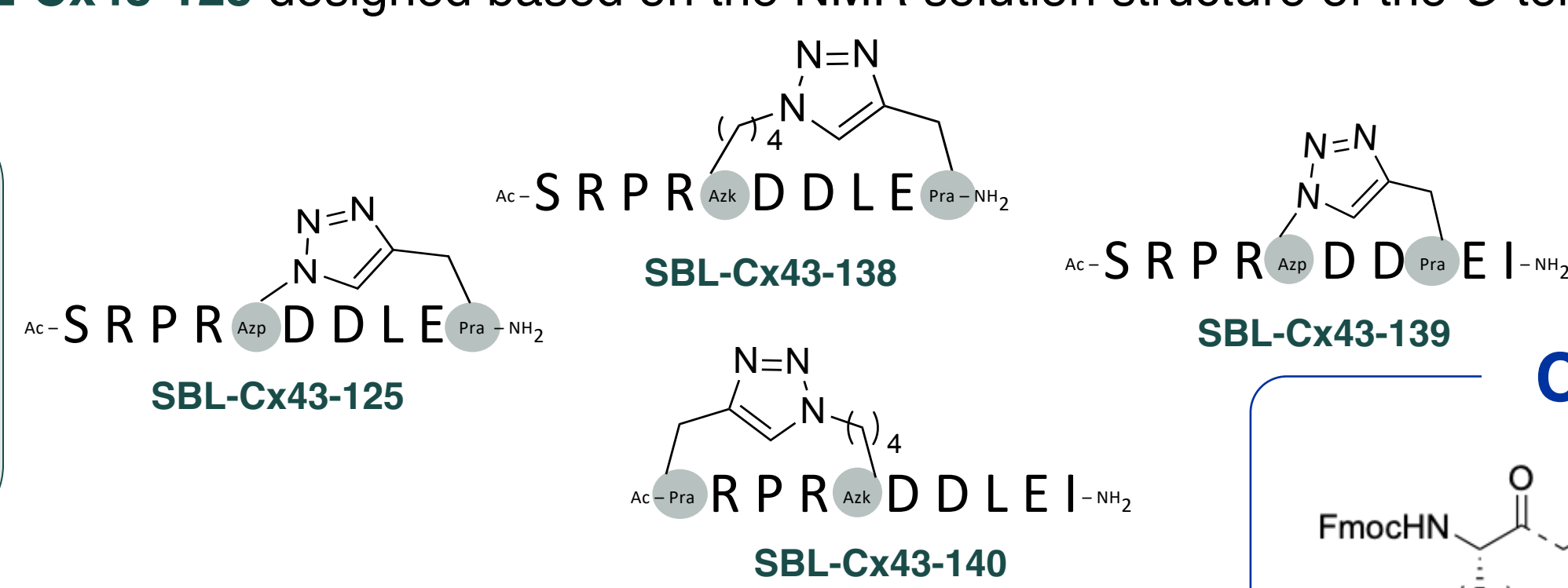
Peptide inhibitors mimicking a segment of the sequences of the proteins, in particular **CT10** and **αCT1** (including a cell-penetrating peptide (CPP)) for Cx43 and **¹⁰Panx1** for Panx1, have shown to be promising **therapeutic agents** both by *in vitro* and *in vivo* experiments.² However, their rapid proteolytic cleavage and low bioavailability make them poor systemically-applied drug candidates.³ To overcome these limitations and fine-tune the properties of the **¹⁰Panx1** and **CT10** peptides, a series of chemical strategies, including **cyclization** and the introduction of **lipidic motifs**, were investigated.^{4,5}

DESIGN AND SYNTHESIS

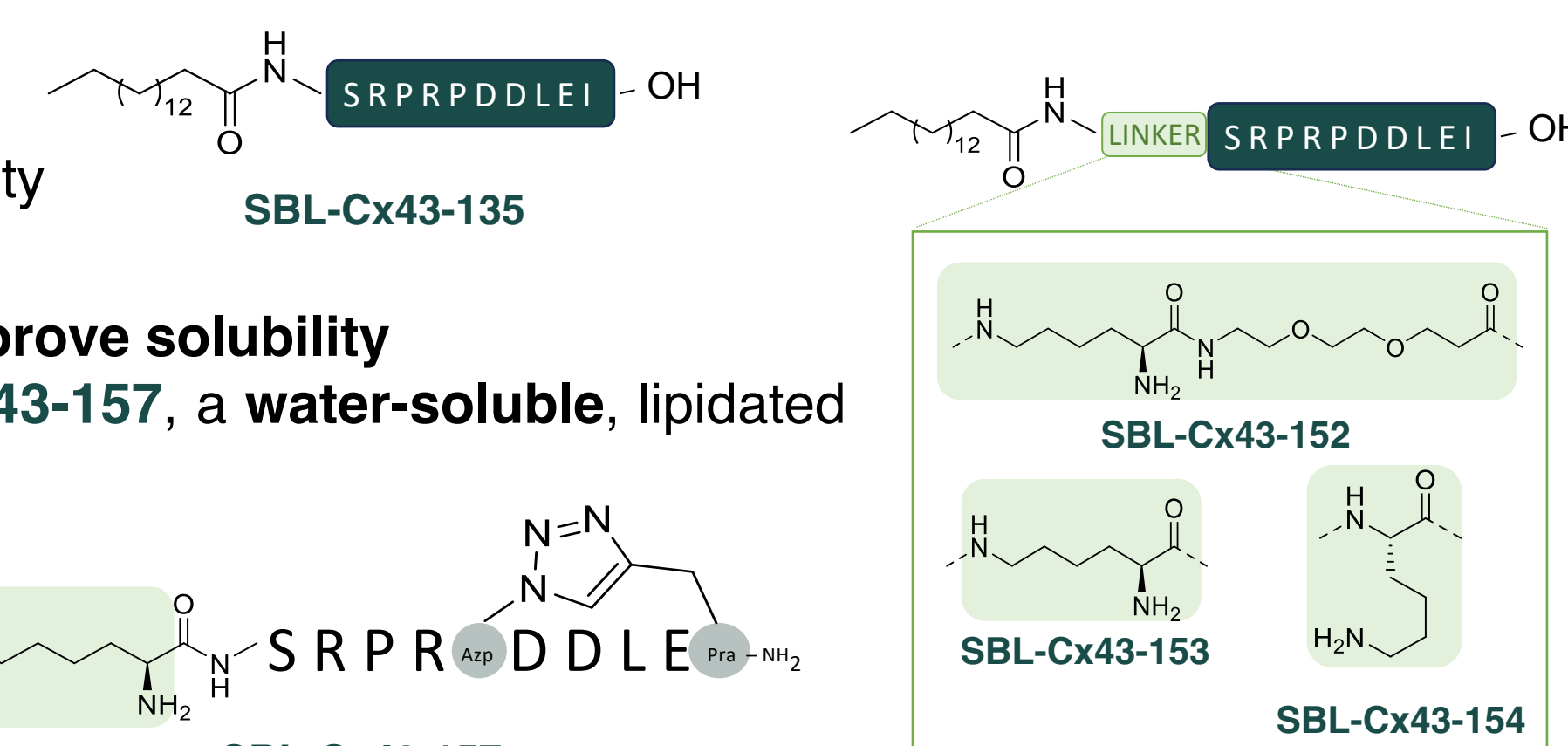
CT10 H-SRPRPDDLEI-OH

- Alanine scan did not pinpoint any residues as essential for biological activity
- Cyclic peptide **SBL-Cx43-125** designed based on the NMR solution structure of the C-terminal domain⁶

Variation of linker position and flexibility allowed for generation of a set of cyclic peptide inhibitors

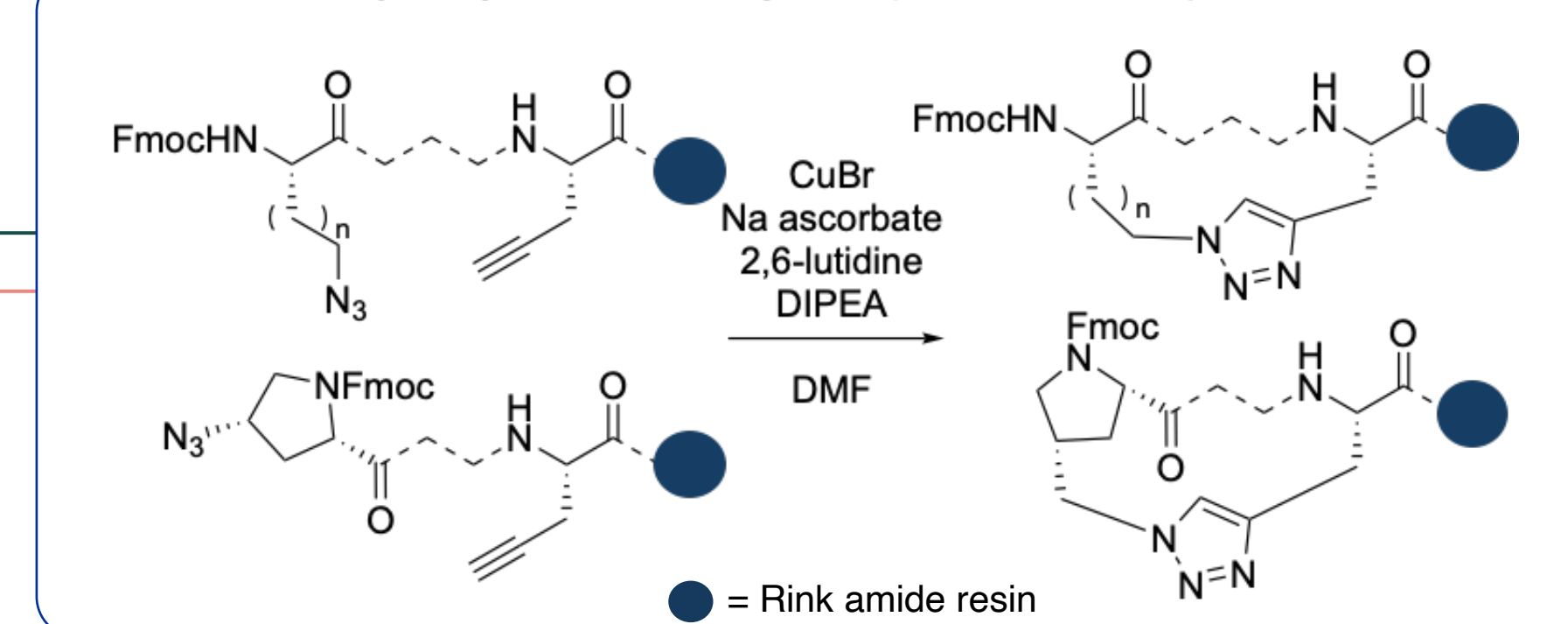


- The use of **lipidic motifs** to improve cell permeability was investigated
- SBL-Cx43-135** shows reduced water solubility



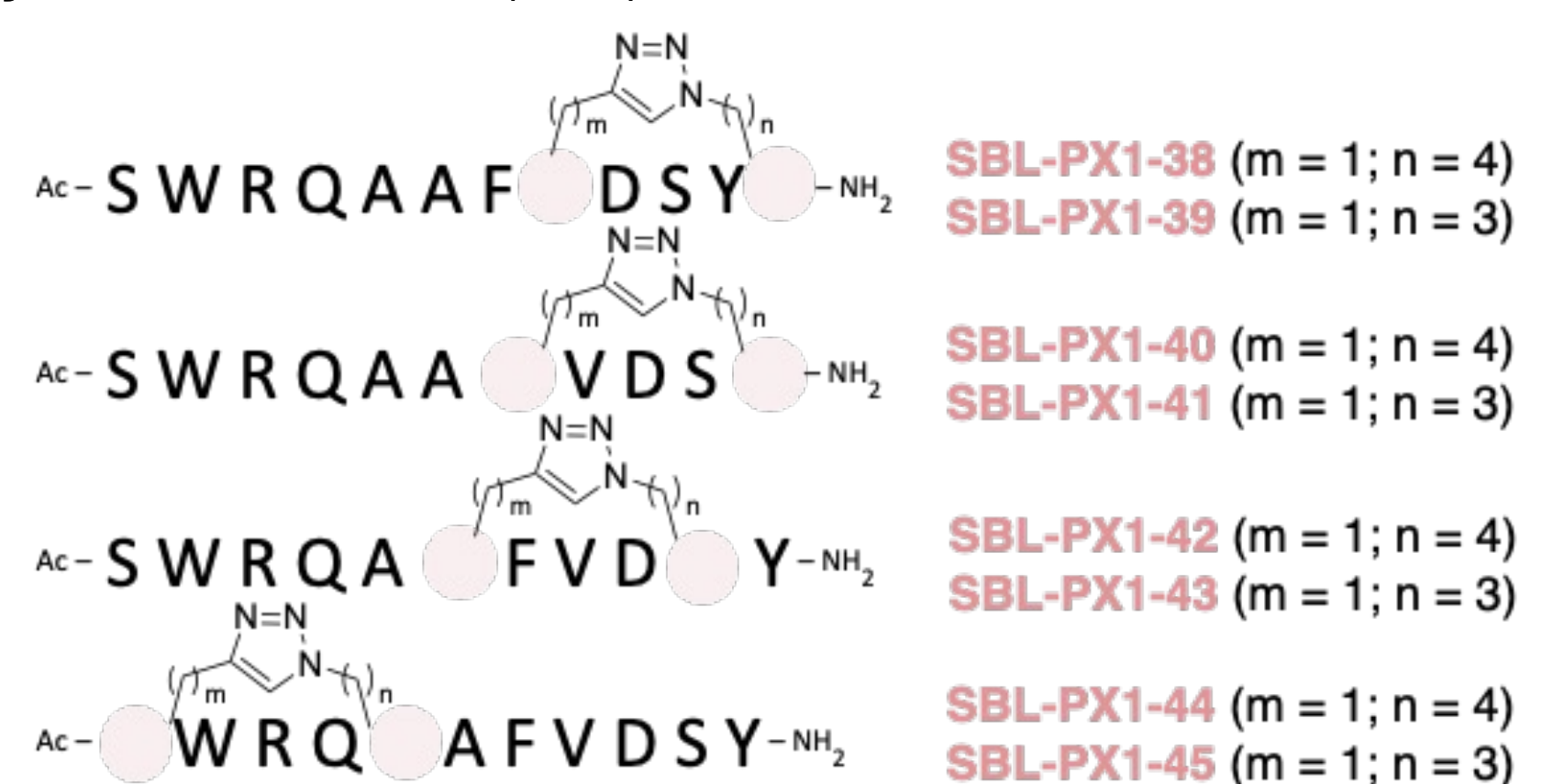
- Different **polar linkers** were introduced to **improve solubility**
- The best linker was used to design **SBL-Cx43-157**, a **water-soluble, lipidated** cyclic peptide based on **SBL-Cx43-125**

CYCLIZATION STRATEGY

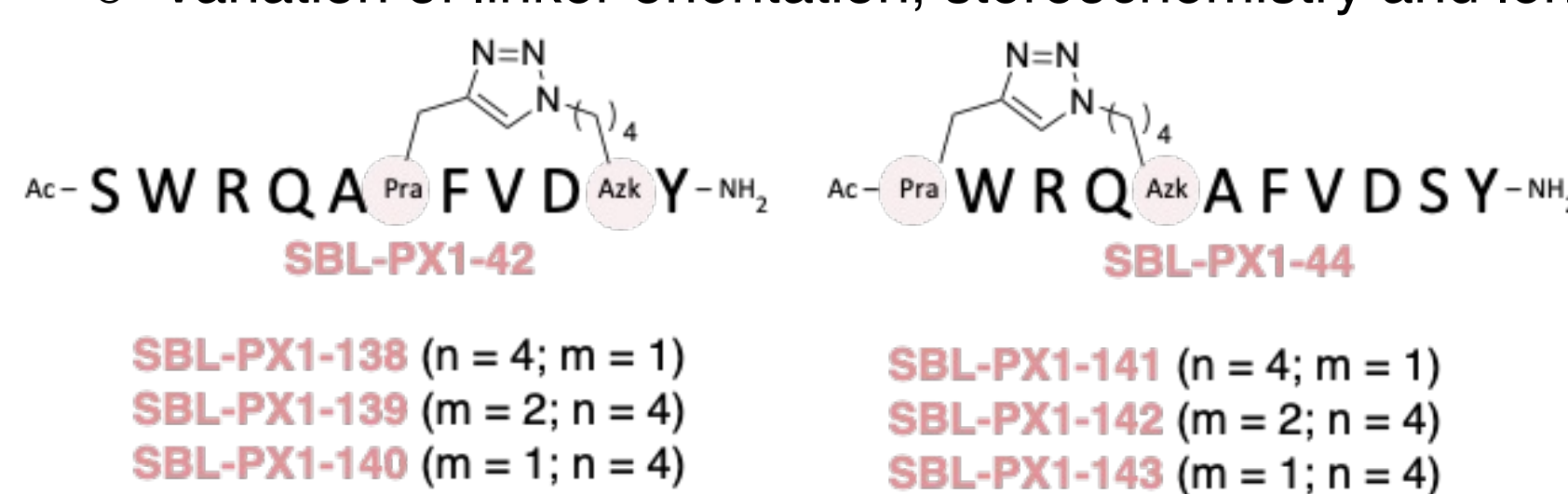


¹⁰Panx1 H-WRQAAFVDSY-OH

- Cyclization scan** at (*i,i+4*) distance to stabilize helical conformation



- Lead peptides **SBL-PX1-42** and **SBL-PX1-44**
- Variation of linker orientation, stereochemistry and length

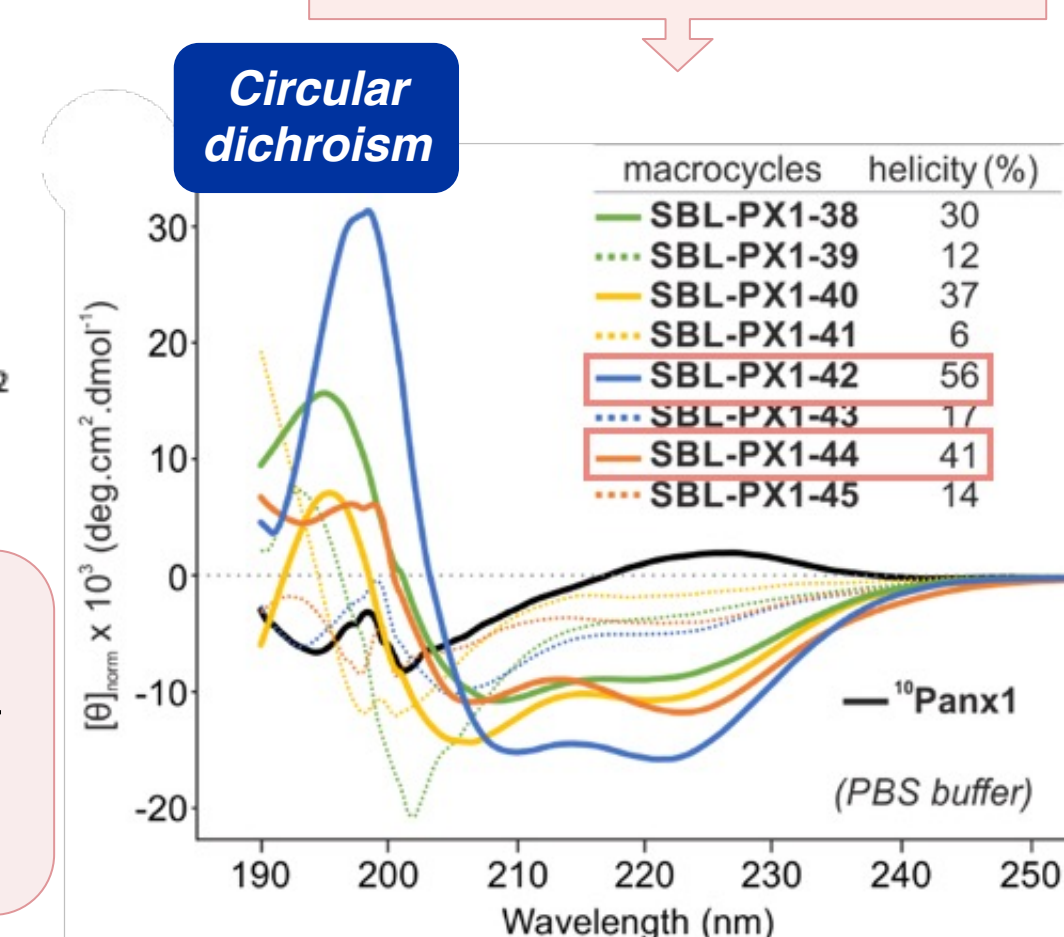


- The structures of lead peptides **SBL-PX1-42** and **SBL-PX1-44** were combined into a **"double stapled"** structure



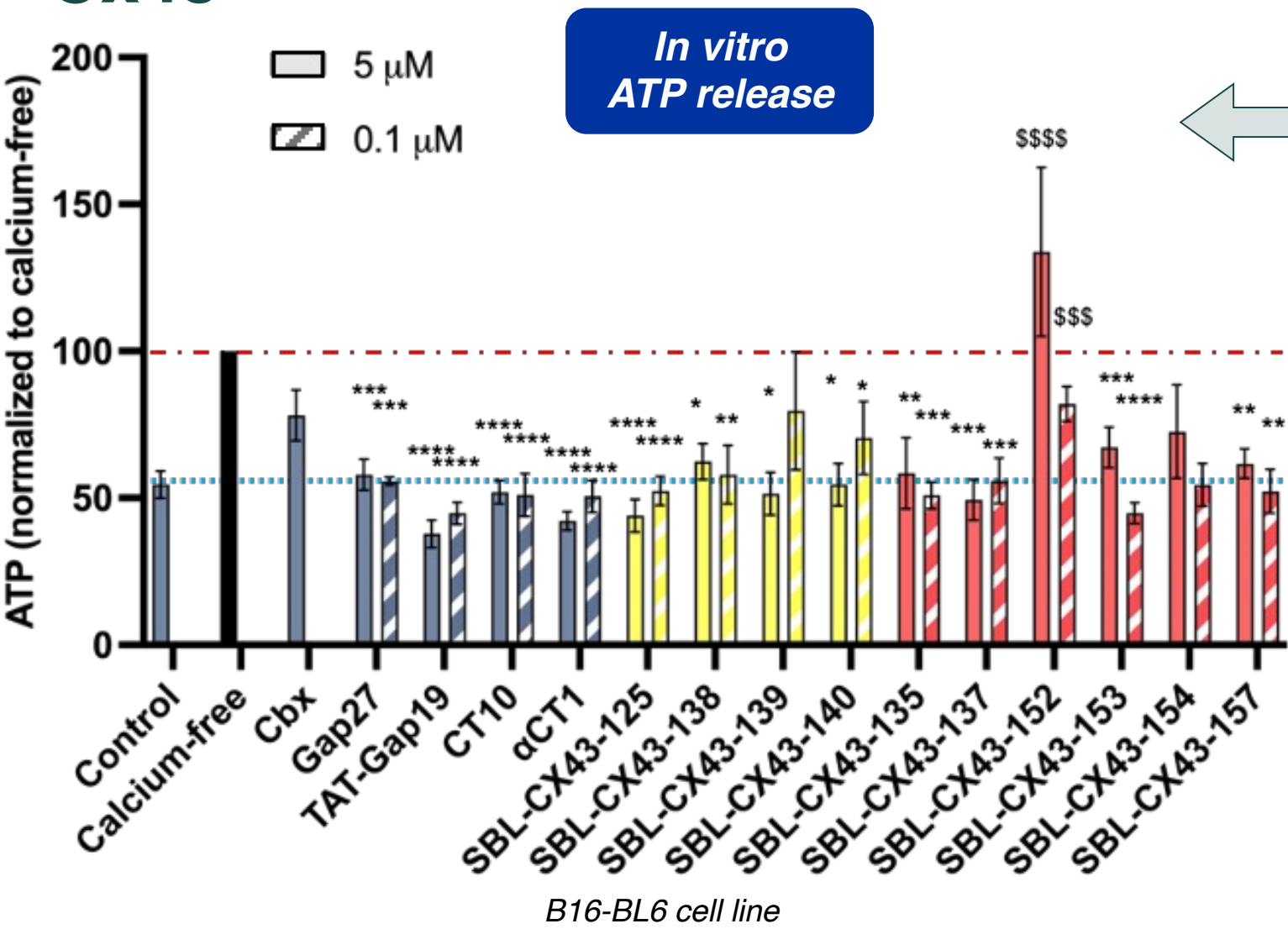
The addition of a **solubilizing tail** (H-K-Ahx) led to the generation of a proteolytically stable, water-soluble bicyclic peptide

The **α-helical secondary structure** is stabilized



BIOLOGICAL ACTIVITY

Cx43

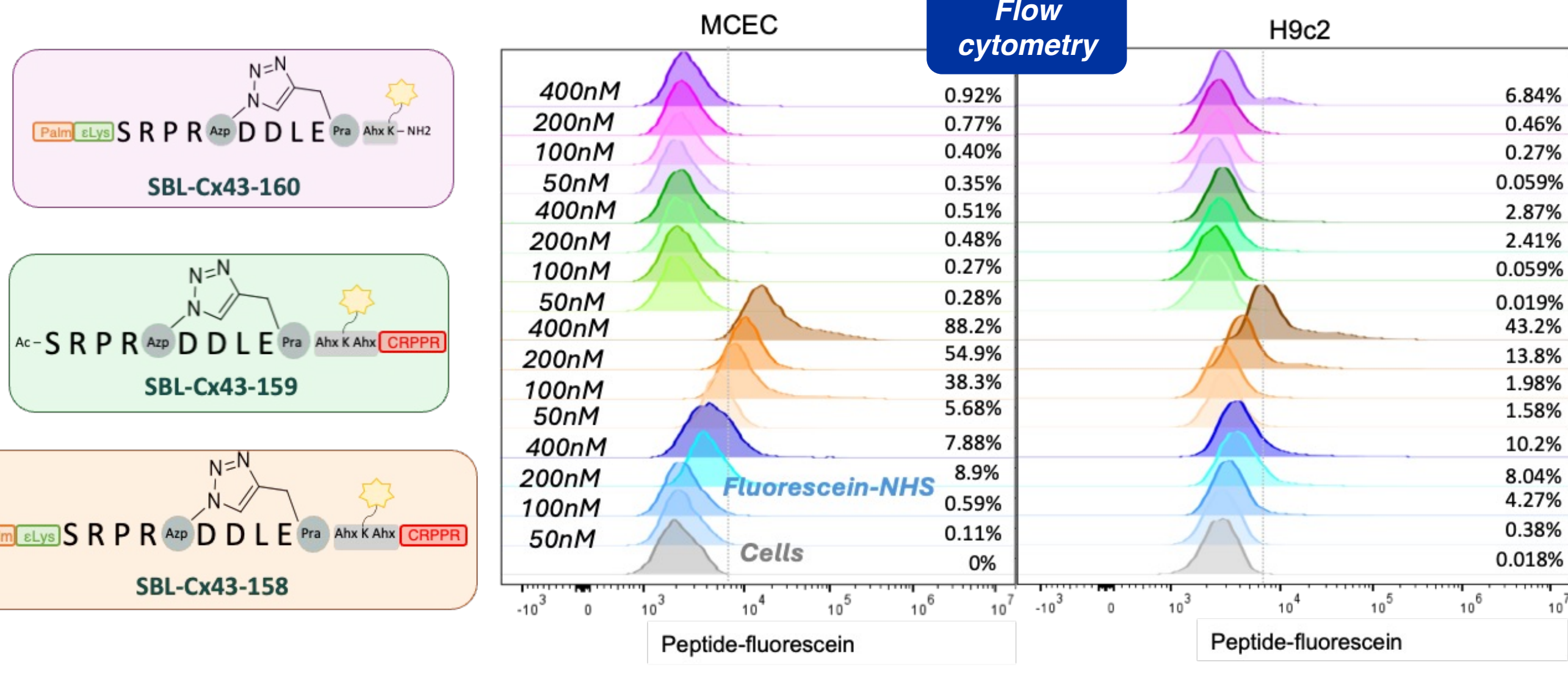


At **5 μM**, all the cyclic peptides have an inhibitory activity comparable to that of **αCT1** without a CPP.

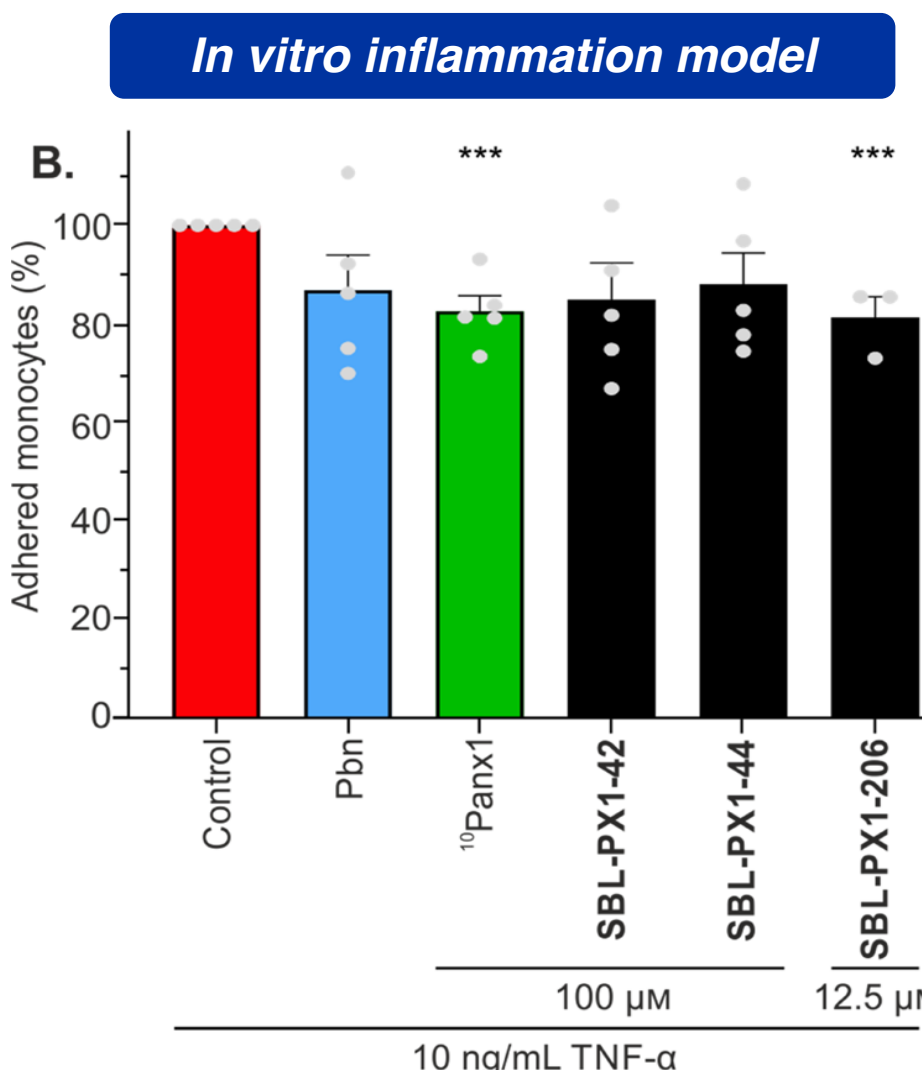
SBL-Cx43-125 retains its activity at a concentration as low as **100 nM**.

The **length** of the linker affects activity of the lipidated peptides **SBL-Cx43-157** shows a higher inhibitory capacity compared to the peptide based on **CT10** (**SBL-Cx43-135**).

The cyclic, lipidated peptide was linked to a **cardiac endothelium-targeting peptide**⁷ and binding was tested by flow cytometry together with two control peptides in different cardiac cell lines. **SBL-Cx43-158** binds preferentially to a mouse cardiac endothelial cell line (MCEC).



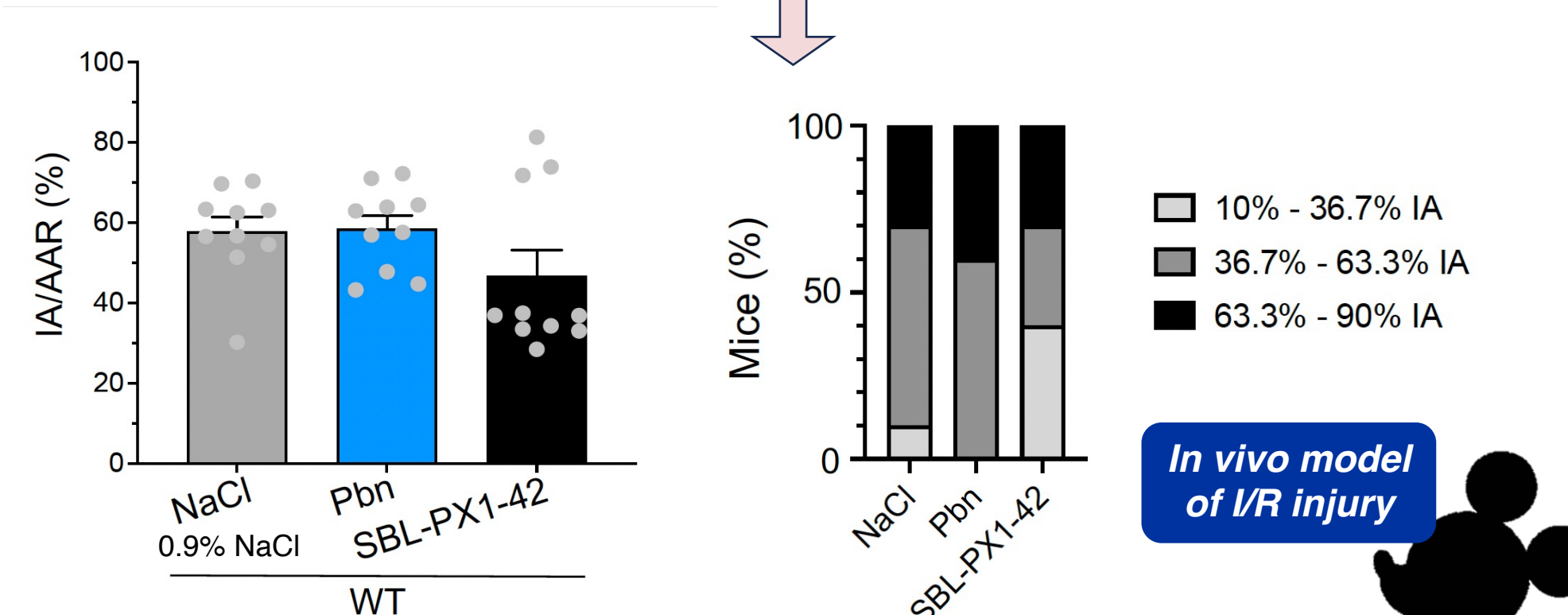
Panx1



SBL-PX1-42 and **SBL-PX1-44** reduce the adhesion of THP-1 monocytes to a TNF-α-activated endothelial monolayer by about 20% at 100 μM, similarly to **¹⁰Panx1**. The same effect was reached by the water-soluble "double stapled" **SBL-PX1-206** at **12.5 μM**.

SBL-PX1-42 was tested in an *in vivo* mouse model of I/R injury. 30 min ischemia followed by 24h reperfusion was induced in WT and Panx1^{-/-} mice by ligation of the left anterior descending coronary artery. 90 μM **SBL-PX1-42** was injected intravenously at 5 min before reperfusion.

SBL-PX1-42 decreased the infarct size by 40% in 7 out of 10 WT mice. No reduction in infarct size was observed in Panx1^{-/-} mice.



PLASMA STABILITY

CT10 $t_{1/2} = 75.86 \pm 4.01$ min

The half-life of **SBL-Cx43-125** largely exceeds 24h

¹⁰Panx1 $t_{1/2} = 2.27 \pm 0.11$ min

SBL-PX1-42 $t_{1/2} = 66.13 \pm 0.52$ min

SBL-PX1-44 $t_{1/2} = 62.42 \pm 2.51$ min

For **SBL-PX1-206**, after 24h, 20% of the intact peptide is still intact in the blood plasma. The only metabolite is the peptide without the solubilizing tail. The stability of the bicyclic structure exceeds 24h.

CONCLUSIONS

- SBL-Cx43-125** is a potent and proteolytically stable inhibitor of Cx43 HC activity *in vitro*, showing an activity comparable to that of **αCT1** at submicromolar concentrations **without requiring a CPP**.

- The designed **solubilizing linkers** allowed to obtain **water-soluble lipidated peptides** of increased lipophilicity, which might be favourable for *in vivo* bioavailability.

- The presence of an endothelial cell-targeting motif in **SBL-Cx43-158** increased the binding of this peptide to MCECs compared to the cardiomyocyte-like cell line H9c2.

- Cyclic peptides **SBL-PX1-42** and **SBL-PX1-44** **reduce monocyte adhesion to the endothelium** (the first step in the inflammatory cascade) by 20%, similarly as **¹⁰Panx1**, with a 30-fold **longer half-life**.

- The combination of the staples in the peptides led to **SBL-PX1-206**. This bicyclic peptide reaches a similar anti-inflammatory activity *in vitro* at **low concentration (12.5 μM)**.

- Preliminary *in vivo* results on **SBL-PX1-42** show its ability to **decrease infarct size by 40%** in 7 out of 10 WT mice in an acute model of **myocardial I/R injury**.

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