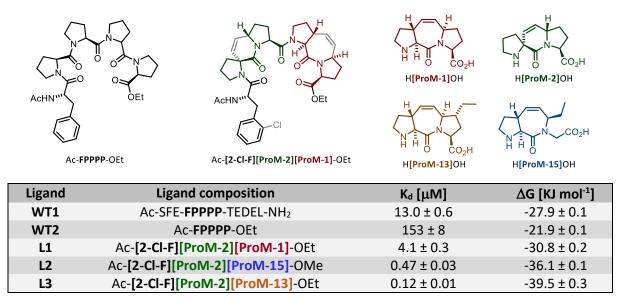
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Designed EVH1 inhibitors derived from synthetic diproline mimetics (ProMs) rigidified in a PPII helix conformation

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We combined computational modeling, X-ray crystallography, organic synthesis and biological investigations to develop small-molecule inhibitors for the Ena/VASP EVH1 domain, which is involved in the migration and chemotaxis (metastasis) of invasive cancer cells.^[1] Using synthetic diproline mimetics such as **ProM-1**,^[2a] **ProM-2**,^[2b] **ProM-13**^[2c] and **ProM-15**^[2d] we succeeded to reduce the wild-type oligopeptide sequence **WT1** to an equivalent of the pentapeptidic proline rich core sequence **WT2**, which itself only shows weak binding. By freezing pairs of prolines in a PPII helix conformation (by formal introduction of a vinylidene bridge) in combination with the introduction of a chlorine atom to the phenylalanine residue, the first nanomolar EVH1 inhibitors (L1-L3) were obtained.^[1]



Besides exhibiting strong affinity, the optimized inhibitors were shown to bind to the EVH1 domain in the canonical fashion. While *in vitro* cell migration experiments proved the inhibition of chemotaxis, the extravasation of MDA-MB-231 cancer cells was demonstrated *in vivo* using transgenic zebrafish embryos.^[1b] Notably, the compounds showed high selectivity towards EVH1 in comparison to other proline-rich motif-recognizing (off-target) protein domains (Fyn SH3, Profilin, Yap1 WW1) as revealed by ¹H¹⁵N HSQC NMR experiments.

References:

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