

Peptide-Vehicle Library

Overview

In the three years from FY 2008 to FY 2010, we promoted Okinawa Innovation Research Consignment Project "Developing cancer therapeutic drugs and test methods using peptide libraries". We designed a peptide-vehicle in which a cell surface recognition peptide, a cell penetrating peptide and an anticancer agent are conjugated as a candidate substance for "cell targeting therapeutic drug", and studied to construct a library. We will provide "Peptide-Vehicle Library" constructed based on the results. In recent years, problems such as QOL decrease due to side effects, patent disappearance of anticancer substances are a problem, and it is expected that drugs will be delivered that anticancer drugs specifically to cancer cells. We hope to use it as a tool of drug discovery research.

Reference Hirata, A., Nokihara, K. Construction of peptide-vehicles, bioconjugates having modules of cancer cell surface capture and cell-penetrating peptide with anticancer agents, *Tetrahedron Lett.* **2014**, 55, 4091-4094.

Peptide No.	Sequence(C terminal amide)	Peptide No.	Sequence(C terminal amide)
HiPeV-001F	aTAMRA-GPKSKRKVYGRKKRRQRRR	HiPeV-005D	Cys(S-EMCA-Gly-CPT)-YGRKKRRQRRRcNNGRGEQc
HiPeV-002F	aTAMRA-GPKSKRKVRQIKIYFQNRRMKWKK	HiPeV-006D	Cys(S-EMCA-Gly-Iri)-YGRKKRRQRRRcNNGRGEQc
HiPeV-003F	aTAMRA-GPKSKRKVRRRRRRRRR	HiPeV-007D	Cys(S-EMCA-Gly-Eto)-YGRKKRRQRRRcNNGRGEQc
HiPeV-004F	aTAMRA-GPKSKRKVRRRRRNTRRRNRRRVR	HiPeV-008D	Cys(S-EMCA-Gly-Pac)-YGRKKRRQRRRcNNGRGEQc
HiPeV-005F	aTAMRA-GKKNQKKRYGRKKRRQRRR	HiPeV-009D	Cys(S-EMCA-Gly-CPT)-RQIKIYFQNRRMKWKKcNNGRGEQc
HiPeV-006F	aTAMRA-GKKNQKKRQRQIKIYFQNRRMKWKK	HiPeV-010D	Cys(S-EMCA-Gly-Iri)-RQIKIYFQNRRMKWKKcNNGRGEQc
HiPeV-007F	aTAMRA-GKKNQKKRRRRRRRRR	HiPeV-011D	Cys(S-EMCA-Gly-Eto)-RQIKIYFQNRRMKWKKcNNGRGEQc
HiPeV-008F	aTAMRA-GKKNQKKRRRRRNTRRRNRRRVR	HiPeV-012D	Cys(S-EMCA-Gly-Pac)-RQIKIYFQNRRMKWKKcNNGRGEQc
HiPeV-009F	aTAMRA-GPKSKRKV	HiPeV-013D	Cys(S-EMCA-Gly-CPT)-RRRRRRRRcNNGRGEQc
HiPeV-010F	aTAMRA-GKKNQKKR	HiPeV-014D	Cys(S-EMCA-Gly-Iri)-RRRRRRRR cNNGRGEQc
HiPeV-011F	aTAMRA-GYGRKKRRQRRR	HiPeV-015D	Cys(S-EMCA-Gly-Eto)-RRRRRRRRcNNGRGEQc
HiPeV-012F	aTAMRA-GRQIKIYFQNRRMKWKK	HiPeV-016D	Cys(S-EMCA-Gly-Pac)-RRRRRRRRcNNGRGEQc
HiPeV-013F	aTAMRA-GRRRRRRRR	HiPeV-017D	Cys(S-EMCA-Gly-CPT)-RRRRNRTRRNRRRVcNNGRGEQc
HiPeV-014F	aTAMRA-GRRRRNRTRRNRRRVR	HiPeV-018D	Cys(S-EMCA-Gly-Iri)-RRRRNRTRRNRRRVcNNGRGEQc
HiPeV-015F	TAMRA-G-YKQC(Acm)HHKGGKKGSG	HiPeV-019D	Cys(S-EMCA-Gly-Eto)-RRRRNRTRRNRRRVcNNGRGEQc
HiPeV-016F	TAMRA-cNGRGEQcYGRKKRRQRRR	HiPeV-020D	Cys(S-EMCA-Gly-Pac)-RRRRNRTRRNRRRVcNNGRGEQc
HiPeV-017F	TAMRA-cNGRGEQcRQIKIYFQNRRMKWKK		
HiPeV-018F	TAMRA-cNGRGEQcRRRRRRRRR		
HiPeV-019F	TAMRA-cNGRGEQcRRRRNRTRRNRRRVR		
HiPeV-020F	TAMRA-YGRKKRRQRRRcNNGRGEQc		
HiPeV-021F	TAMRA-RQIKIYFQNRRMKWKKcNNGRGEQc		
HiPeV-022F	TAMRA-RRRRRRRRcNNGRGEQc		
HiPeV-023F	TAMRA-RRRRNRTRRNRRRVcNNGRGEQc		
HiPeV-024F	FAM-GYGRKKRRQRRR		
HiPeV-025F	FAM-GRQIKIYFQNRRMKWKK		
HiPeV-026F	FAM-GRRRRRRRR		
HiPeV-001D	C(S-EMCA-Gly-CPT)-YGRKKRRQRRR		
HiPeV-002D	C(S-EMCA-Gly-Iri)-YGRKKRRQRRR		
HiPeV-003D	C(S-EMCA-Gly-Eto)-YGRKKRRQRRR		
HiPeV-004D	C(S-EMCA-Gly-Pac)-YGRKKRRQRRR		

