

peptide: 33

.(TMT6plex)GK(TMT6plex)VK(TMT6plex)VGVNGFGR_0
(TMT6plex)GK(TMT6plex)VK(TMT6plex)VGVNGFGR_2
AAFNNGSK(TMT6plex)_0
AGAHLQGGAK(TMT6plex)_0
AGAHLQGGAK(TMT6plex)_2
AGAHLQGGAK(TMT6plex)R_0
FHGTVK(TMT6plex)_0
GALQNIIPAS(TMT6plex)TGAAG(TMT6plex)AVGK(TMT6plex)_0
GALQNIIPASTGAAG(TMT6plex)_2
GALQNIIPASTGAAG(TMT6plex)_3

GILGYTEHQVVS(TMT6plex)SDFNS(TMT6plex)DTHSSTFDAGAGIALNDHFVK(TMT6plex)_0
GILGYTEHQVVS(TMT6plex)SDFNSDTHSSTFDAGAGIALNDHFVK(TMT6plex)_0
GILGYTEHQVVS(TMT6plex)DFNS(TMT6plex)DTHSSTFDAGAGIALNDHFVK(TMT6plex)_0
GILGYTEHQVVS(TMT6plex)DFNSDTHSSTFDAGAGIALNDHFVK(TMT6plex)_0
IIS(TMT6plex)NASC(Carbamidomethyl)TTNC(Carbamidomethyl)LAPLAK(TMT6plex)_0
IK(TMT6plex)WGDAGAEYVVS(TMT6plex)TGVFTTMEK(TMT6plex)AGAHLQGGAK(TMT6plex)R_0
LEK(TMT6plex)PAK(TMT6plex)YDDIK(TMT6plex)_0
LEK(TMT6plex)WGDAGAEYVVS(TMT6plex)TGVFTTMEK(TMT6plex)_3
LEK(TMT6plex)PAK(TMT6plex)YDDIK(TMT6plex)_3
LEK(TMT6plex)PAK(TMT6plex)YDDIK(TMT6plex)K(TMT6plex)_0
LIS(TMT6plex)WYDNEFGYSNR_0

OAS(TMT6plex)EGPLK(TMT6plex)_0
VIHDMFVIGELMTTVVHAITATQK(TMT6plex)_0
VIIS(TMT6plex)APS(TMT6plex)ADAPMFVGMVNHKEK(TMT6plex)_0
VIIS(TMT6plex)APSADAPM(Oxidation)FVMGVNHKEK(TMT6plex)_0
VIIS(TMT6plex)APSADAPMFVM(Oxidation)GVNHKEK(TMT6plex)_0
VIIS(TMT6plex)APSADAPMFVM(Oxidation)GVNHKEK(TMT6plex)_3
VIIS(TMT6plex)APSADAPMFVGMVNHKEK(TMT6plex)_0
VIISADAPM(Oxidation)FVMGVNHKEK(TMT6plex)_0
VIISADAPMFVGMVNHKEK(TMT6plex)_0
VIISADAPMFVGMVNHKEK(TMT6plex)_2

VIISAPSADAPMFVGMVNHKEK(TMT6plex)YDNSLK(TMT6plex)_0
VPTANVS(TMT6plex)VVDLTC(Carbamidomethyl)R_0
VPTANVS(TMT6plex)VVDLTC(Carbamidomethyl)R_2
VPTANVS(TMT6plex)VVDLTC(Carbamidomethyl)R_3
VVDLMAHM(Oxidation)ASK(TMT6plex)E_0
VVDLMAHMAS(TMT6plex)K(TMT6plex)E_0
VVDLMAHMASK(TMT6plex)_0
VVDLMAHMASK(TMT6plex)_3
VVDLMAHMASK(TMT6plex)E_0
VVDLMAHMASK(TMT6plex)E_2

WGDAGAEYVVS(TMT6plex)TGVFTTMEK(TMT6plex)_0
WGDAGAEYVVS(TMT6plex)TGVFTTMEK(TMT6plex)_2
WGDAGAEYVVS(TMT6plex)TGVFTTMEK(TMT6plex)_3
WGDAGAEYVVS(TMT6plex)TGVFTTMEK(TMT6plex)R_0
YDNSLK(TMT6plex)_0
YDNSLK(TMT6plex)_2

