

Supporting Information

Unambiguous Identification of Serine and Threonine Pyrophosphorylation Using Neutral-Loss-Triggered Electron-Transfer/Higher-Energy Collision Dissociation (EThcD)

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Representative HCD MS/MS spectra

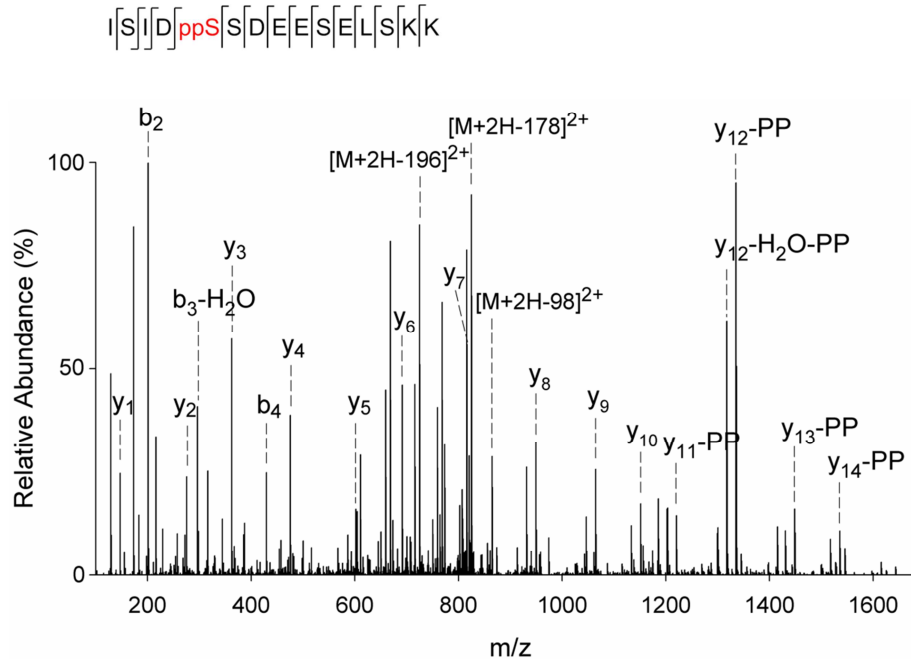


Figure S1. HCD MS/MS spectrum of doubly charged peptide ISIDppSSDEESELKK (**PP-4**)

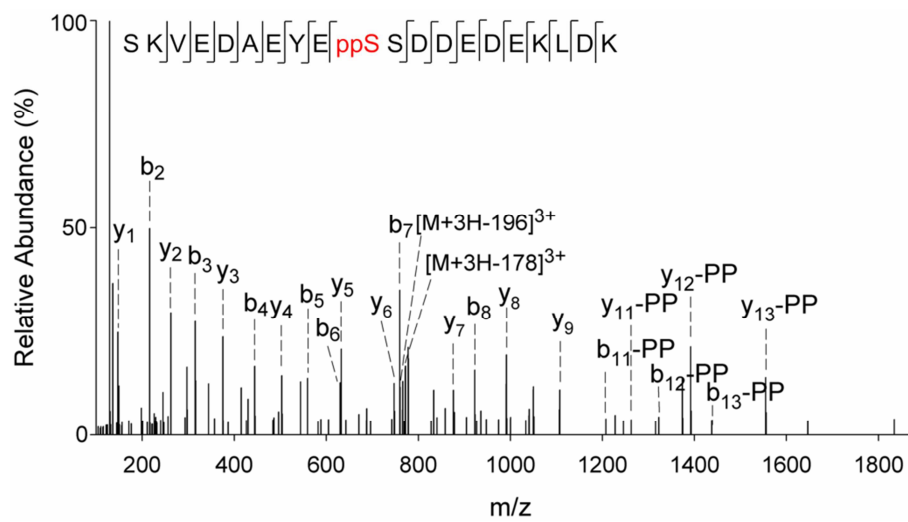


Figure S2. HCD MS/MS spectrum of triply charged peptide SKVEDAEYEppSSDDEDEKLDDK (**PP-6**)

Representative CID MS/MS spectra

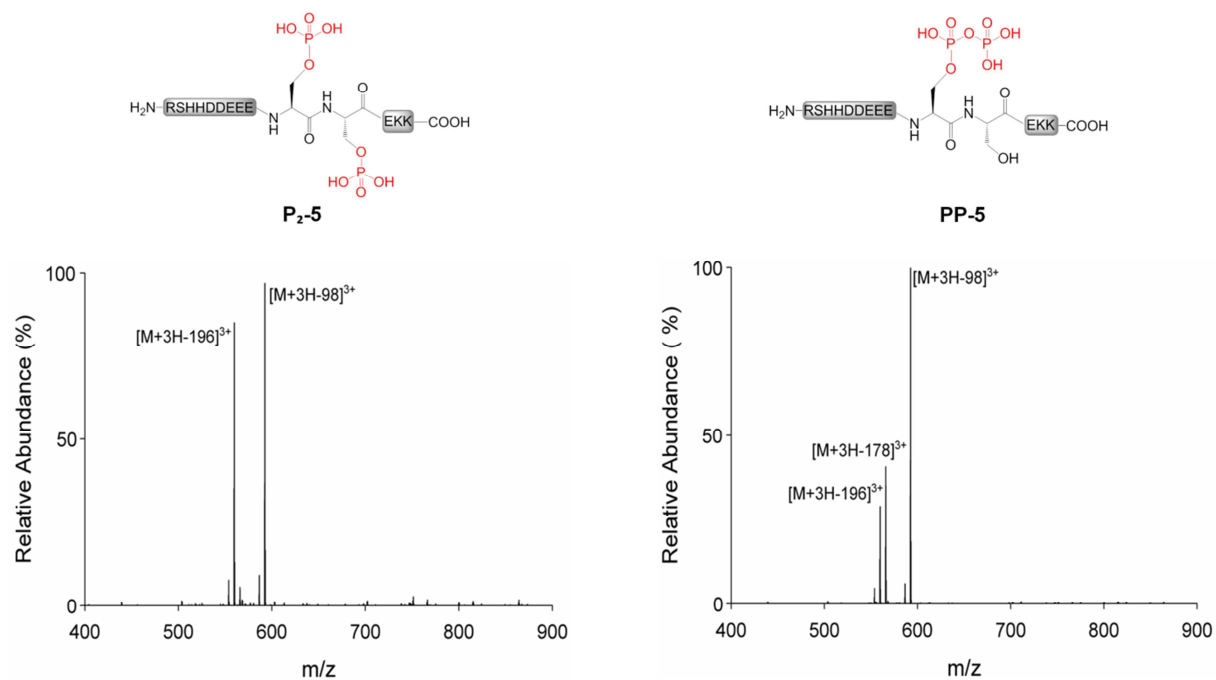


Figure S3. Comparison of CID MS/MS spectrum of triply charged peptide RSHHDDEEEpSpSEKK (**P₂-5**) and peptide RSHHDDEEEppSSEKK (**PP-5**)

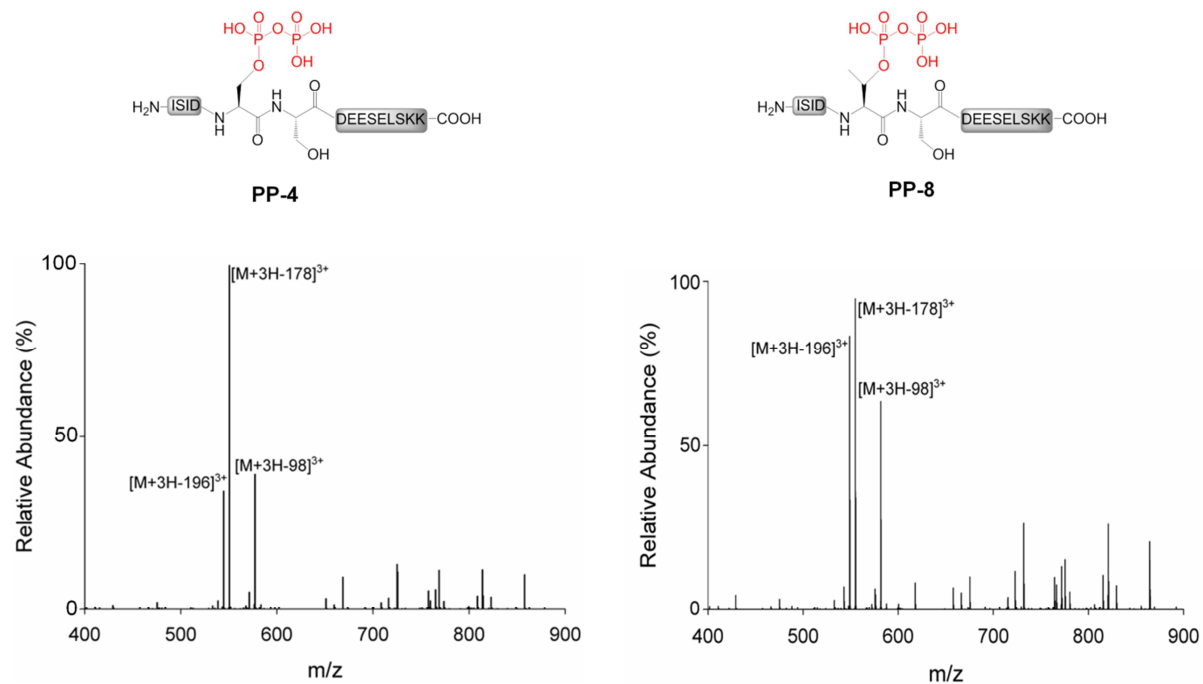


Figure S4. Comparison of CID MS/MS spectrum of triply charged peptide ISIDppSSDEESELKK (**PP-4**) and peptide ISIDppTSDEESELKK (**PP-8**)

Representative EThcD MS/MS spectra of triply charged precursor ions

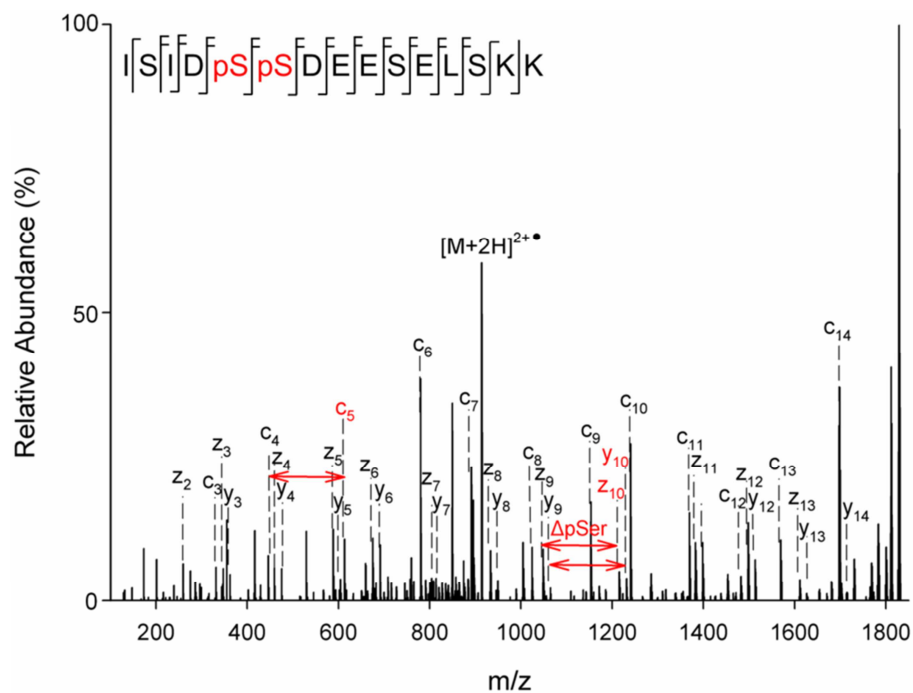


Figure S5. EThcD MS/MS spectrum of triply charged peptide ISIDpSpSDEESEL SKK (P₂-4)

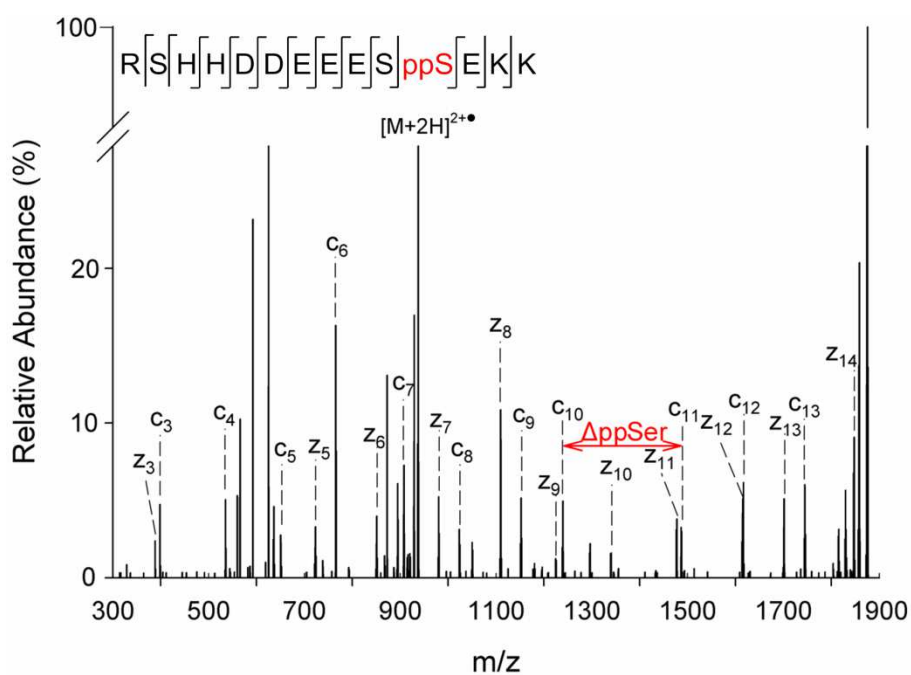


Figure S6. EThcD MS/MS spectrum of triply charged peptide RSHHDDEEESppSEKK (PP-5)

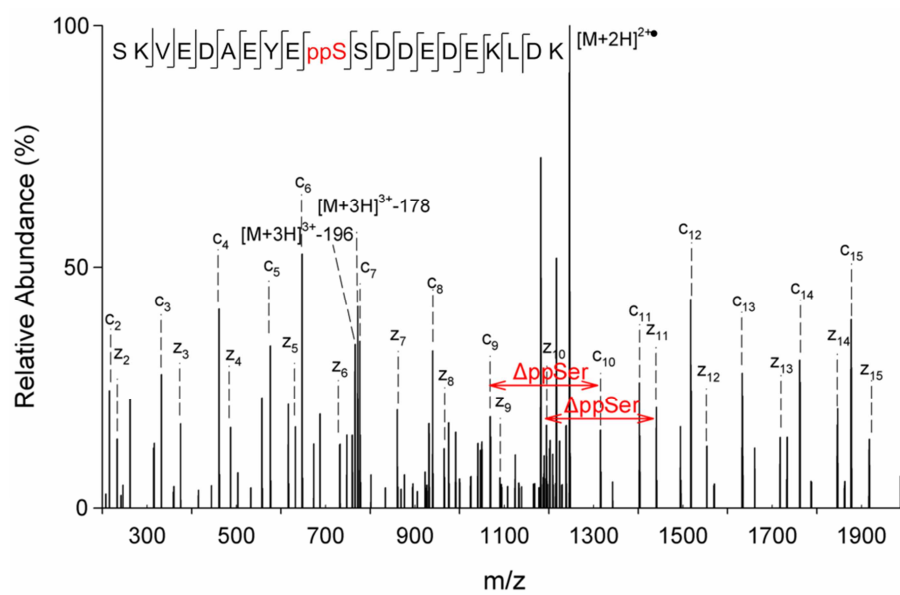


Figure S7. ETHcD MS/MS spectrum of triply charged peptide SKVEDAEYEppSSDDEDEKLDK (PP-6)

Representative EThcD MS/MS spectra of doubly charged precursor ions

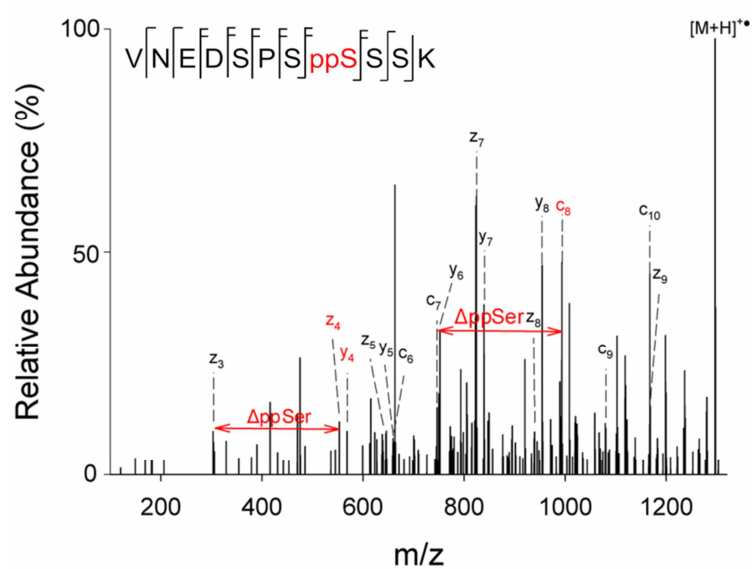


Figure S8. EThcD MS/MS spectrum of doubly charged peptide VNEDSPppSSSK (**PP-1**)

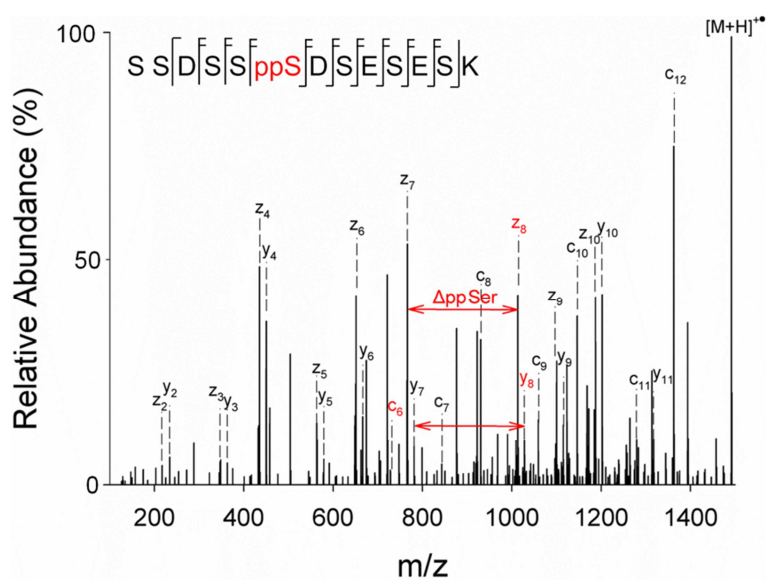


Figure S9. EThcD MS/MS spectrum of doubly charged peptide SSDSppSDSESESK (**PP-3**)

Chromatographic behavior of di- and pyrophosphorylated peptides

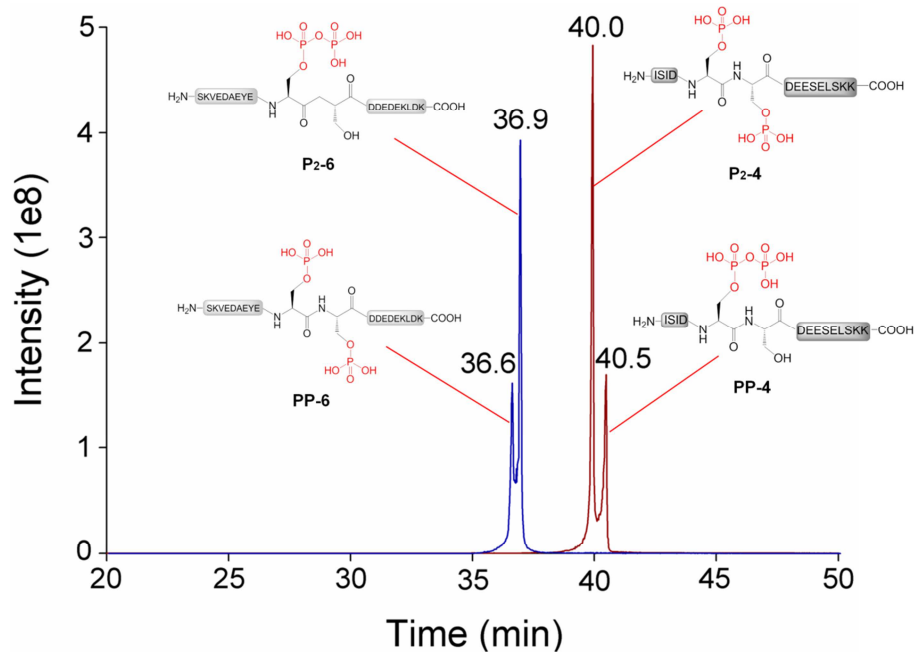


Figure S10. Extracted ion chromatograms (XIC) of the synthetic peptides **P2-6**, **PP-6** (m/z 830.982; blue line), **P2-4** and **PP-4** (m/z 913.873; red line) showing the similar chromatographic behavior of di- and pyrophosphorylated peptides. The doubly phosphorylated peptides exhibit shorter retention times than the pyrophosphorylated counterpart.

Spike-in experiment peptide ISIDppTSDEESELKK (PP-8)

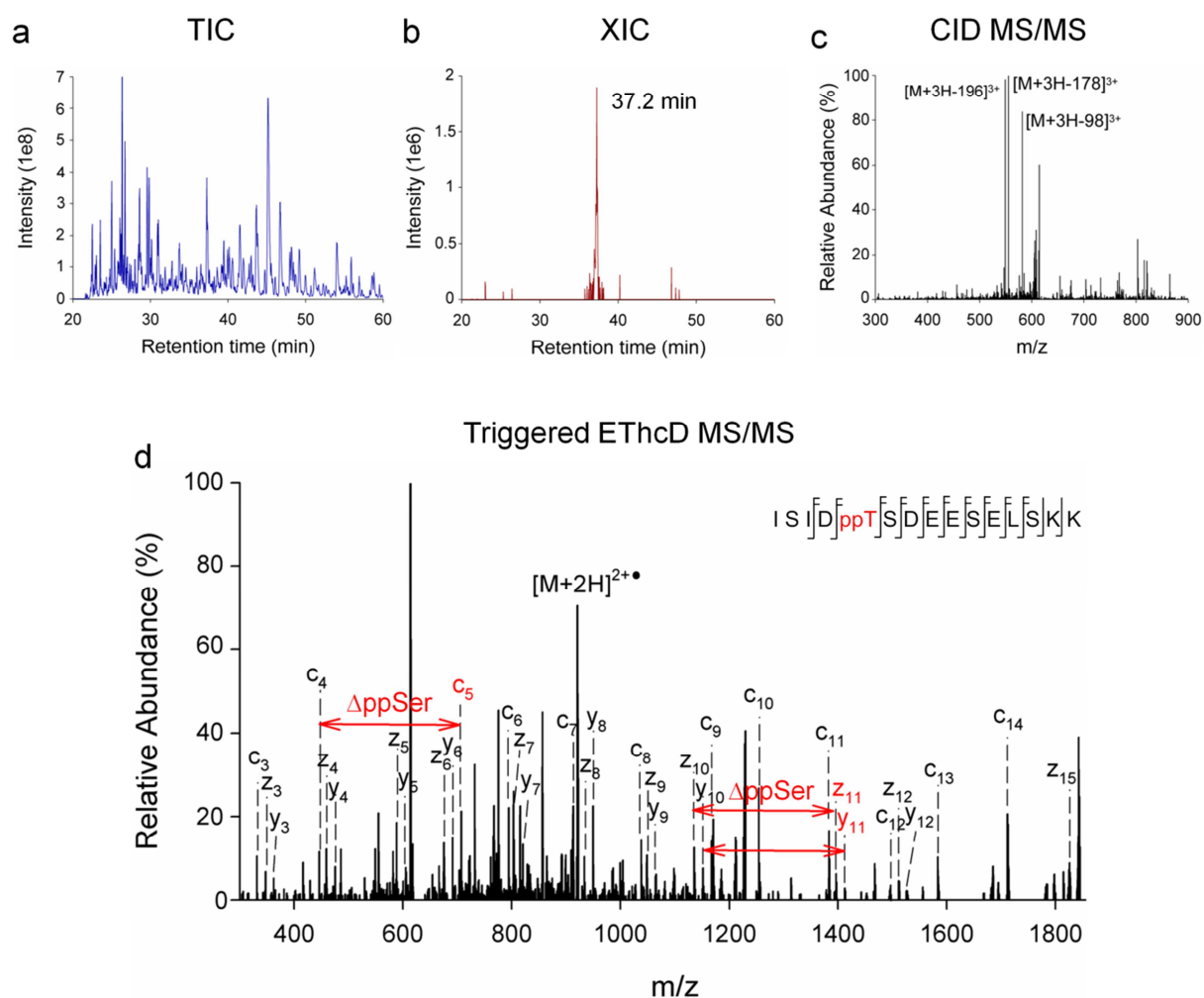
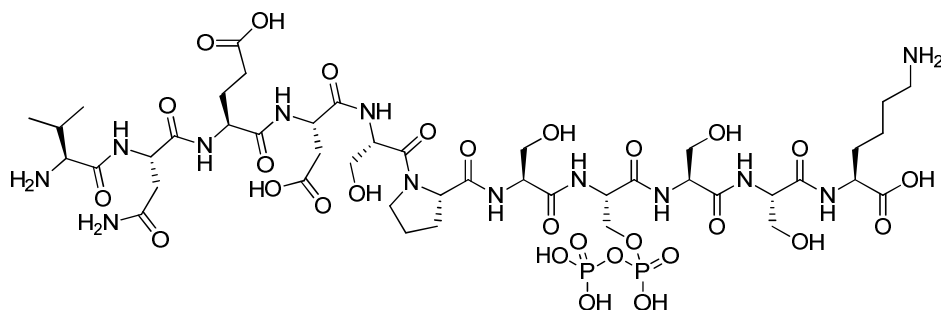


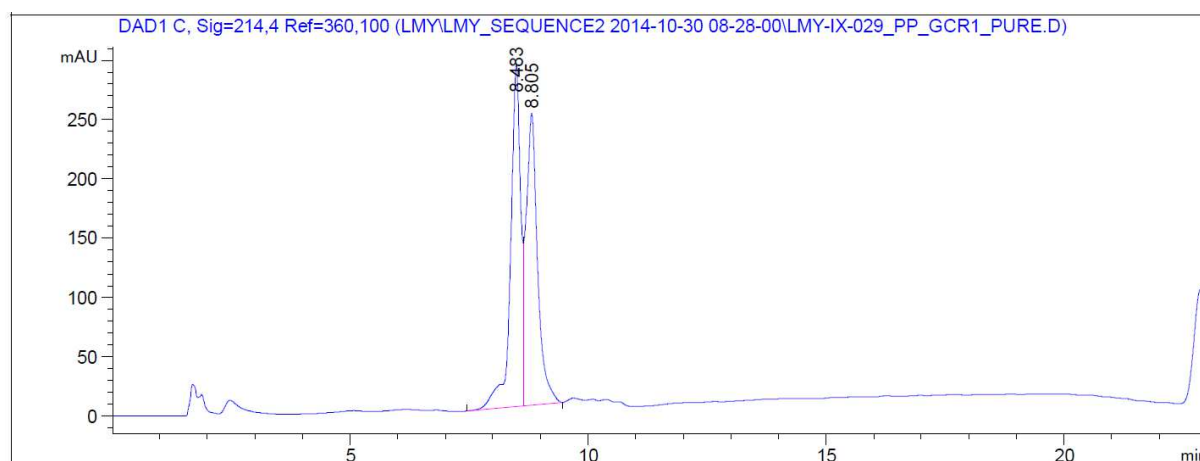
Figure S11. Detection of a synthetic pyrophosphopeptide in a spike-in experiment using the DDNL-ETHcD approach. (a) Total ion chromatogram (TIC) of the HeLa protein digest. (b) Extracted ion chromatogram (XIC) m/z 614.255 of the synthetic peptide **PP-8**. (c) CID MS/MS spectrum of peptide **PP-8** acquired at a retention time of 37.23 min indicating dominant neutral losses of 98, 178 and 196 Da. (d) Triggered ETHcD MS/MS spectrum of the pyrophosphorylated peptide ISIDppTSDEESELKK showing gapless sequence coverage without loss of the labile modification. Fragment ions pinpointing the site of modification are labeled in red.

Characterization of synthetic pyrophosphopeptides

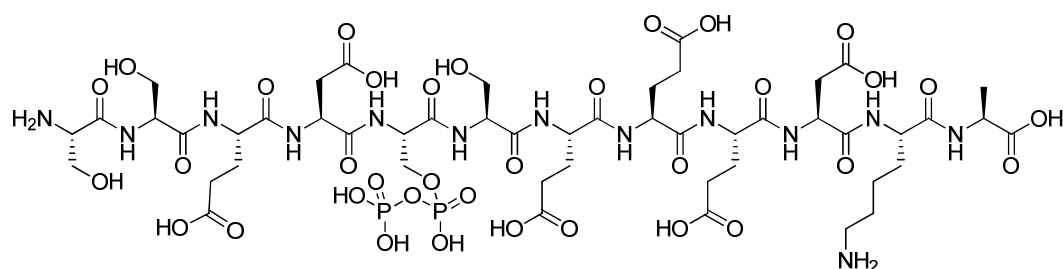
Peptide **PP-1** (VNEDSPSpSSSK)



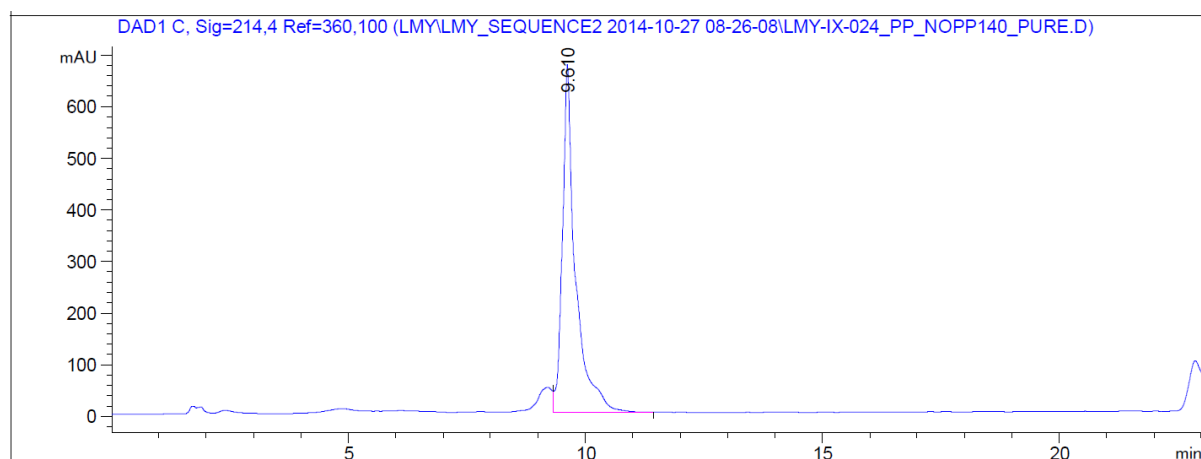
The purity of the isolated peptide was confirmed by analytical HPLC [C18; $t = 0$ min 0% of solvent B in solvent A, $t = 20$ min 20% of solvent B in solvent A; 1 mL/min; 214 nm; $T_R = 8.483$ min & $T_R = 8.805$ min]. Both peaks in the analytical HPLC were collected and HRMS confirmed that both fractions contained pyrophosphopeptide **PP-1** as the primary product (HRMS (Bulk Material) $[M+H]^+$ calcd for $C_{44}H_{76}N_{13}O_{28}P_2^+$ 1296.4393, found 1296.4463; $[M+2H]^{2+}$ calcd for $C_{44}H_{77}N_{13}O_{28}P_2^{2+}$ 648.7233, found 648.7255; $[M+2H+Na]^{3+}$ calcd for $C_{44}H_{77}N_{13}NaO_{28}P_2^{3+}$ 440.8133, found 440.8070) and ^{31}P NMR (202 MHz, H_2O , pH = 7.70) δ -6.78 (d, $^2J_{P-P} = 27.8$ Hz), -10.73 (d, $^2J_{P-P} = 19.9$ Hz).



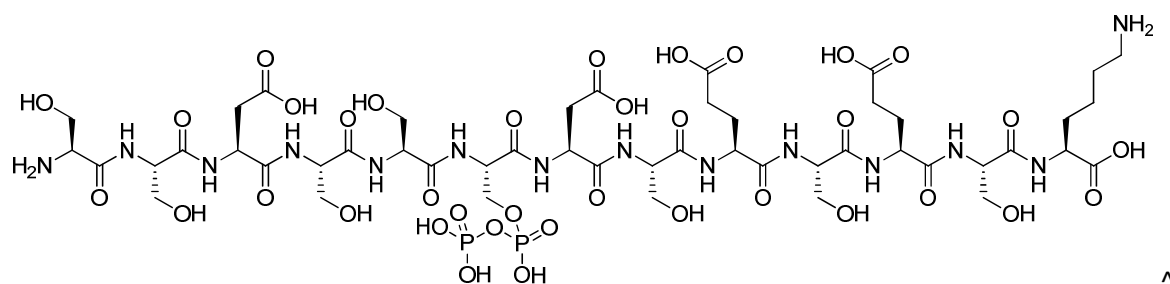
Peptide **PP-2** (SSEDppSSEEDKA)



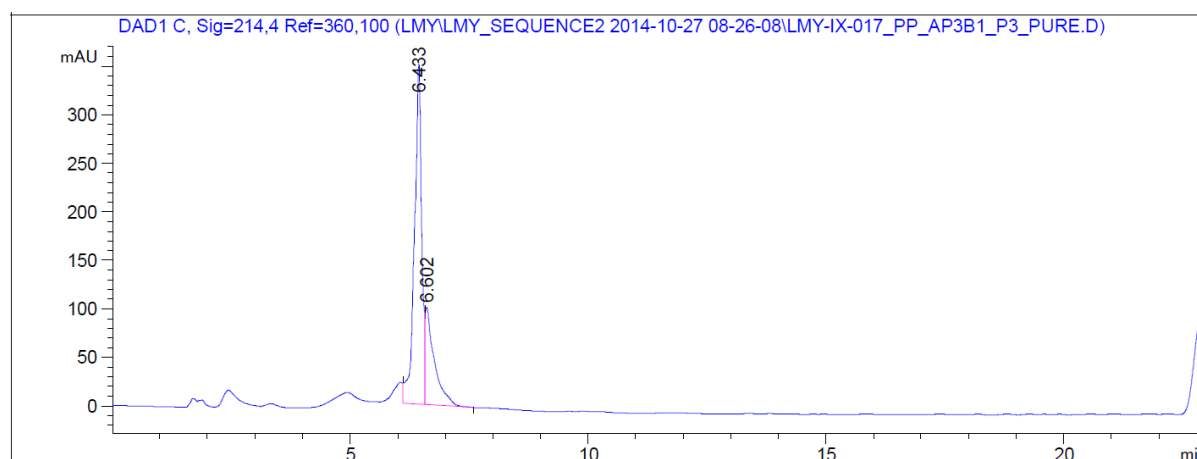
The purity of the isolated peptide was confirmed by analytical HPLC [C18; $t = 0$ min 0% of solvent B in solvent A, $t = 20$ min 20% of solvent B in solvent A; 1 mL/min; 214 nm; $T_R = 9.610$ min], and the identity of pyrophosphopeptide **PP-2** was confirmed by mass spectrometry (HRMS $[M+H]^+$ calcd for $C_{49}H_{80}N_{13}O_{35}P_2^+$ 1472.4350, found 1472.4332; $[M+2H]^{2+}$ calcd for $C_{49}H_{81}N_{13}O_{35}P_2^{2+}$ 736.7211, found 736.72125; $[M+2H+Na]^{3+}$ calcd for $C_{49}H_{81}N_{13}NaO_{35}P_2^{3+}$ 499.4794, found 499.4697) and ^{31}P NMR (202 MHz, H_2O , pH = 7.59) δ -6.84 (d, $^2J_{P-P} = 20.5$ Hz), -10.87 (d, $^2J_{P-P} = 21.1$ Hz).



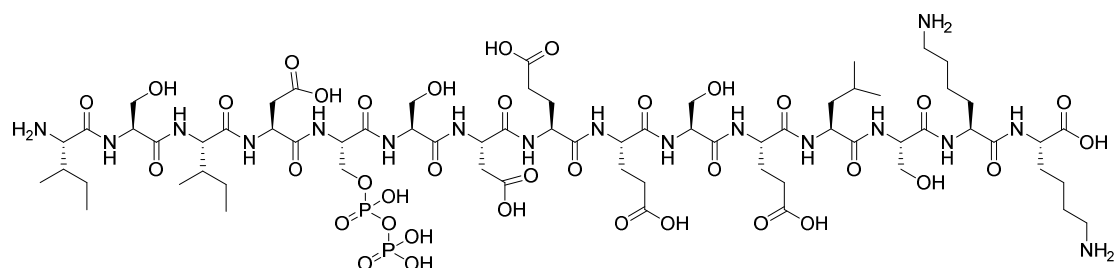
Peptide **PP-3** (SSEDppSSEEDKA)



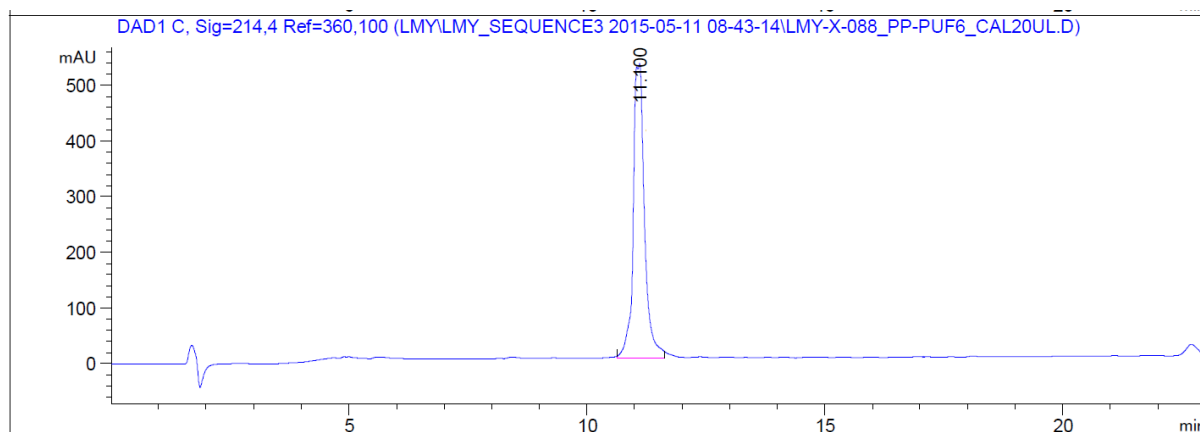
The purity of the isolated peptide was confirmed by analytical HPLC [C18; t = 0 min 0% of solvent B in solvent A, t = 20 min 20% of solvent B in solvent A; 1 mL/min; 214 nm; T_R = 6.433 min], and the identity of the pyrophosphopeptide **PP-3** was confirmed by mass spectrometry (HRMS [M+H+Na]²⁺ calcd for C₄₈H₈₁N₁₄O₃₆P₂⁺ 1491.4408, found 1491.4410; [M+2H]²⁺ calcd for C₄₈H₈₂N₁₄O₃₆P₂²⁺ 746.2240, found 746.2255; [M+2H+Na]³⁺ calcd for C₄₈H₈₂N₁₄NaO₃₆P₂³⁺ 505.8147, found 505.8057) and ³¹P NMR (202 MHz, H₂O, pH = 7.67) δ -6.57 (d, ²J_{P-P} 21.8 Hz), -10.81 (d, ²J_{P-P} = 21.1 Hz).



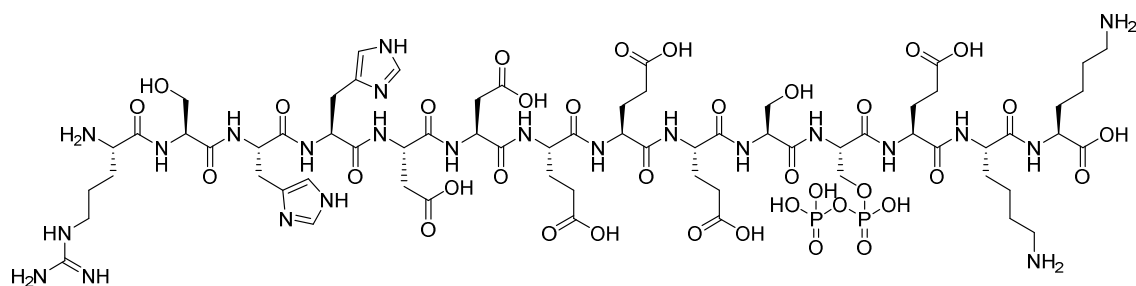
Peptide **PP-4** (ISIDppSSDEESELKK)



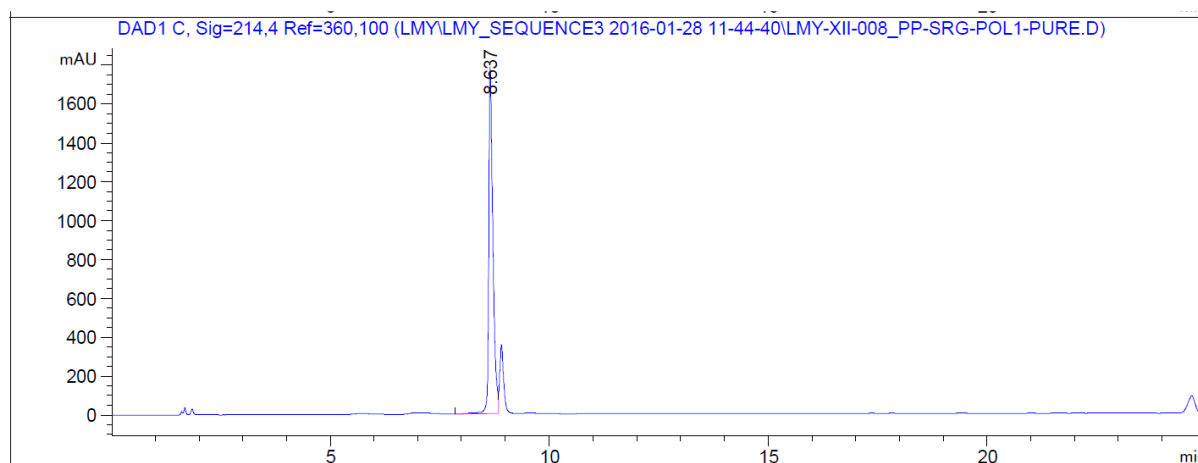
The purity of the isolated peptide was confirmed by analytical HPLC [C18; t = 0 min 0% of solvent B in solvent A; t = 20 min 47% of solvent B in solvent A; 1 mL/min; 214 nm; T_R = 13.957 min], and the identity of pyrophosphopeptide **PP-4** was confirmed by mass spectrometry (HRMS $[M+2H]^{2+}$ calcd for $C_{68}H_{119}N_{17}O_{37}P_2^{2+}$ 913.8709, found 913.8686; $[M+3H]^{3+}$ calcd for $C_{68}H_{119}N_{17}O_{37}P_2^{2+}$ 609.5830, found 609.5823) and ^{31}P NMR (202 MHz, H_2O , pH = 8.07) δ -6.77 (d, $^2J_{P-P}$ = 20.5 Hz), -10.99 (d, $^2J_{P-P}$ = 21.7 Hz).



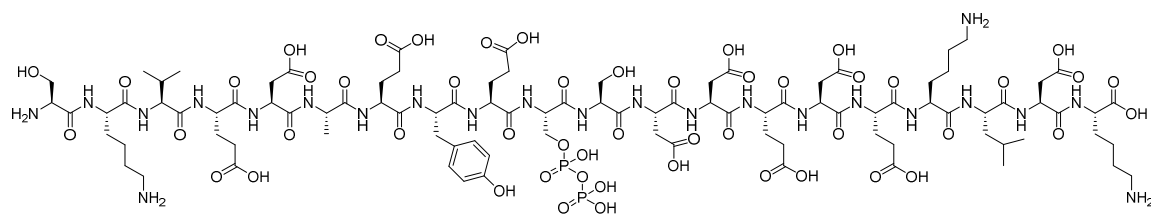
Peptide **PP-5** (RSHHDDEEESppSEKK)



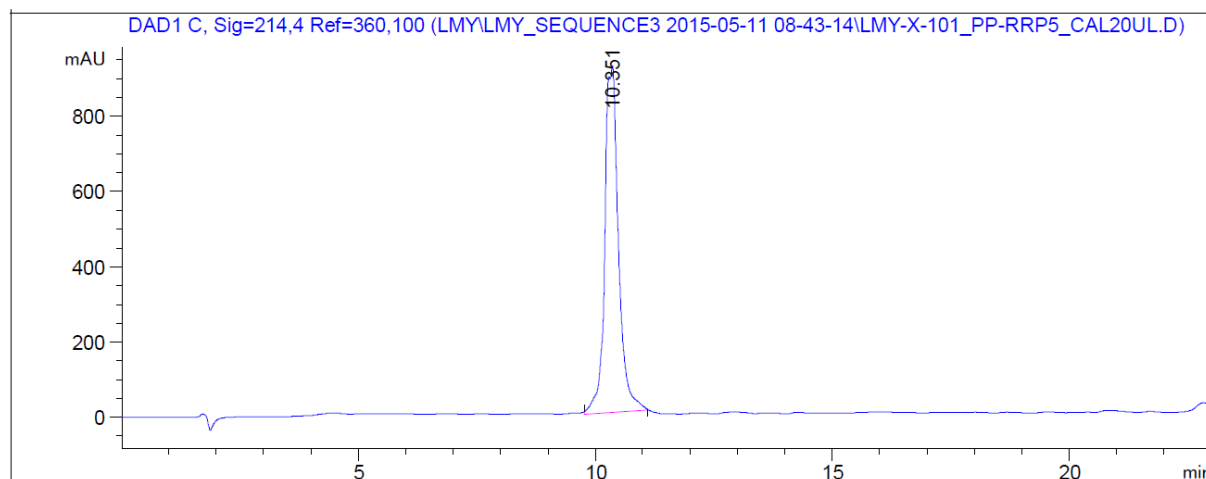
The purity of the isolated peptide was confirmed by analytical HPLC [C18; t = 0 min 3% of solvent B in solvent A; t = 13 min 15% of solvent B in solvent A; 1 mL/min; 214 nm; T_R = 8.637 min], and the identity of pyrophosphopeptide **PP-5** was confirmed by mass spectrometry (HRMS [M+2H]²⁺ calcd for C₉₅H₁₅₁N₂₃O₅₁P₂²⁺ 936.8362, found 936.8294; [M+3H]³⁺ calcd for C₉₅H₁₅₂N₂₃O₅₁P₂³⁺ 624.8908, found 624.8833; [M+Na+3H]⁴⁺ calcd for C₉₅H₁₅₂N₂₃NaO₅₁P₂⁴⁺ 468.9181, found 468.9133) and ³¹P NMR (202 MHz, H₂O, pH = 7.91) δ -6.27 (d, , ²J_{P,P} = 21.8 Hz), -10.76 (d, , ²J_{P,P} = 21.4 Hz).



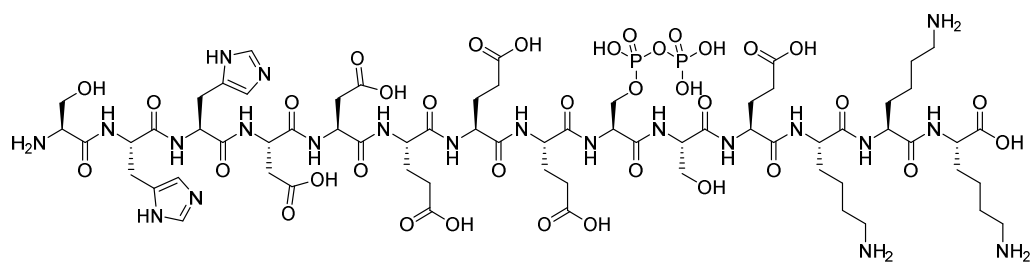
Peptide **PP-6** (SKVEDAEYEppSSDDEDEKLDK)



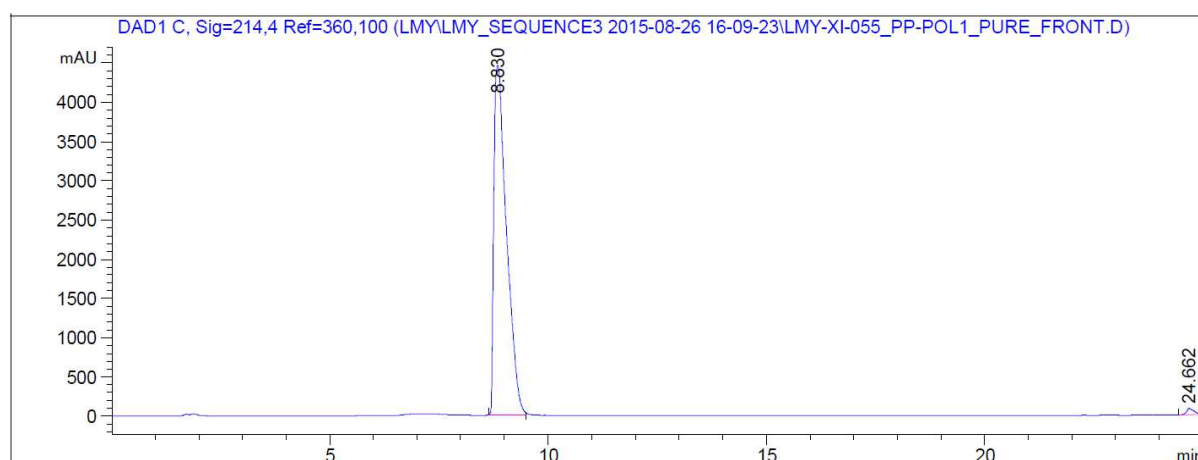
The purity of the isolated peptide was confirmed by analytical HPLC [C18; t = 0 min 0% of solvent B in solvent A; t = 20 min 47% of solvent B in solvent A; 1 mL/min; 214 nm; T_R = 10.351 min], and the identity of pyrophosphopeptide **PP-6** was confirmed by mass spectrometry (HRMS $[M+2H]^{2+}$ calcd for $C_{95}H_{151}N_{23}O_{51}P_2^{2+}$ 1246.4714, found 1246.4687; $[M+3H]^{3+}$ calcd for $C_{95}H_{152}N_{23}O_{51}P_2^{3+}$ 831.3167, found 831.3180; $[M+Na+3H]^{4+}$ calcd for $C_{95}H_{152}N_{23}NaO_{51}P_2^{4+}$ 629.2348, found 629.7307) and ^{31}P NMR (202 MHz, H_2O , pH = 7.94) δ -6.64 (d, , $^2J_{P-P}$ = 21.7 Hz), -10.78 (d, , $^2J_{P-P}$ = 21.5 Hz).



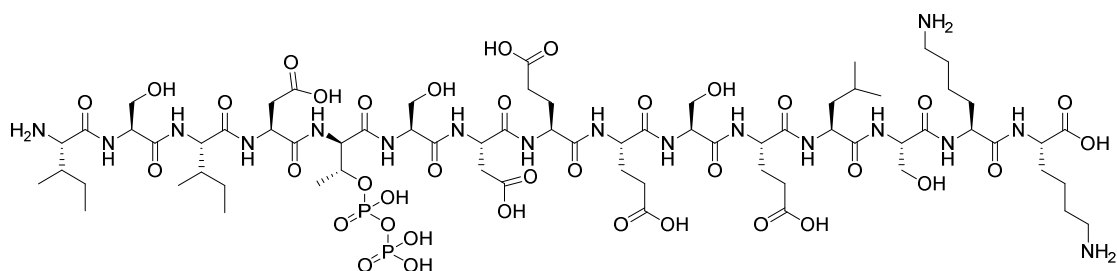
Peptide **PP-7** (SHHDDEEESppSEKKK)



Using Procedure B, pyrophosphopeptide **xxx** was prepared from phosphopeptide **xxx** (11.5 mg, 6.52 μmol) in 2 steps as previously described⁵ to give the title peptide in 28% yield (3.40 mg, 1.84 μmol) over two steps as a white solid. The purity of the isolated peptide was determined by analytical HPLC [C18; $t = 0$ min 0% of solvent B in solvent A; $t = 20$ min 30% of solvent B in solvent A; 1 mL/min; 214 nm; $T_R = 8.830$ min], and the identity of pyrophosphopeptide **PP-7** was confirmed by mass spectrometry (HRMS $[M+2H]^{2+}$ calcd for $C_{67}H_{109}N_{21}O_{36}P_2^{2+}$ 922.8404, found 922.8409; $[M+3H]^{3+}$ calcd for $C_{67}H_{110}N_{21}O_{36}P_2^{3+}$ 615.5627, found 615.5619; $[M+4H]^{4+}$ calcd for $C_{67}H_{111}N_{21}O_{36}P_2^{4+}$ 461.9238, found 461.9221) and ^{31}P NMR (202 MHz, H_2O , pH = 7.75) δ -6.43 (d, $^2J_{P,P} = 21.5$ Hz), -10.86 (d, $^2J_{P,P} = 21.5$ Hz).



Peptide **PP-8** (ISIDppTSDEESELKK)



The purity of the isolated peptide was confirmed by analytical HPLC [C18; $t = 0$ min 0% of solvent B in solvent A; $t = 20$ min 47% of solvent B in solvent A; 1 mL/min; 214 nm; $T_R = 10.463$ min], and the identity of pyrophosphopeptide **PP-8** was confirmed by mass spectrometry (HRMS $[M+2H]^{2+}$ calcd for $C_{69}H_{121}N_{17}O_{37}P_2^{2+}$ 921.8820, found 921.0145; $[M+3H]^{3+}$ calcd for $C_{68}H_{119}N_{17}O_{37}P_2^{2+}$ 614.2549, found 614.3337) and ^{31}P NMR (202 MHz, H_2O , pH = 7.91) δ -6.97 (d, $^2J_{P-P} = 20.4$ Hz), -11.73 (d, $^2J_{P-P} = 20.5$ Hz).

