

Figure 1 displays 15 panels showing the results of 15 MS runs. Each panel displays a network graph with nodes and edges, colored by the number of MS runs (0.125, 0.5, 0.667, 1). The panels are arranged in a grid, with the first row containing 10 panels and the second row containing 5 panels. The x-axis is labeled 'MS runs'.

P06576

peptide: 39

- [K].aHGGYSVFAGVGEr.[T]_3

[K].gFQQILAGEYDHLPEQAFYmVGPIEEAVak.[A]_3

[K].gFQQILAGEYDHLPEQAFYmVGPIEEAVak.[A]_3

[K].gFQQILAGEYDHLPEQAFYmVGPIEEAVak.[A]_4

[K].gFQQILAGEYDHLPEQAFYmVGPIEEAVak.[A]_4

[K].gFQQILAGEYDHLPEQAFYmVGPIEEAVak.[A]_5

[K].gSITSVQAIYVPADDLTDPAPATTFAHLDATTVLSr.[A]_3

[K].gSITSVQAIYVPADDLTDPAPATTFAHLDATTVLSr.[A]_4

[K].iGLFGGAGVGk.[T]_2

[K].iGLFGGAGVGk.[T]_3

[K].iPVGPETLGr.[I]_2

[K].kGSITSVQAIYVPADDLTDPAPATTFAHLDATTVLSr.[A]_4
- [K].qFAPIHAAEPEFMEMSVEQEILVTGik.[V]_3
- [K].qFAPIHAAEPEFMEMSVEQEILVTGik.[V]_4
- [K].qFAPIHAAEPEFMEMSVEQEILVTGik.[V]_4
- [K].sLQDIIAILGMDDELSEEDk.[L]_2
- [K].sLQDIIAILGMDDELSEEDk.[L]_3
- [K].sLQDIIAILGMDDELSEEDk.[L]_3
- [K].sLQDIIAILGMDDELSEEDk.[L]_4
- [K].sLQDIIAILGMDDELSEEDk.[L]_4
- [K].tVLIMELINNvak.[A]_2
- [K].tVLImELINNvak.[A]_3
- [K].tVLIMELINNvak.[A]_3
- [K].vALVYGQmNEPPGAR.[A]_2

[K].vALVYQGMNEPPGAR.[A]_2

[K].vALVYQmNEPPGAR.[A]_3

[K].vALVYQGMNEPPGAR.[A]_3

[K].vLDSGAPIk.[I]_2

[K].vVDLLAPYak.[G]_3

[R].aIAELGIYPavDPLDSTsr.[I]_2

[R].aIAELGIYPavDPLDSTsr.[I]_3

[R].aIAELGIYPavDPLDSTsr.[I]_3

[R].dQEGQDVLLFIDNIFr.[F]_2

[R].dQEGQDVLLFIDNIFr.[F]_3

[R].dQEGQDVLLFIDNIFr.[F]_3

[R].eGNDLYHEmIESGVINLk.[D]_3

[R].eGNDLYHEmIESGVINLk.[D]_3

[R].eGNDLYHEmIESGVINLk.[D]_4

[R].fLSQPFQVAEVFTGHMGk.[L]_3

[R].fLSQPFQVAEVFTGHMGk.[L]_3

[R].fLSDGAPIk.[I]_2

[R].ftQAGSEVSALLGr.[I]_2

[R].ftQAGSEVSALLGr.[I]_3

[R].IMDPNIVGSEHYDVAr.[G]_3

[R].imNVIGEPIDeR.[G]_2

[R].imNVIGEPIDeR.[G]_2

[R].imNVIGEPIDeRGPIk.[T]_3

[R].iPSAVGYQPTLATDMGMTQeR.[I]_3

[R].iPSAVGYQPTLATDMGMTQeR.[I]_3

[R].iPSAVGYQPTLATDMGMTQeR.[I]_3

[R].iPSAVGYQPTLATDMGMTQeR.[I]_3

[R].iVLEVAQHLGESTvr.[T]_3

[R].iAmDGTEGLVr.[G]_2

[R].iAmDGTEGLVr.[G]_2

[R].trEGNDLYHEmIESGVINLk.[D]_3

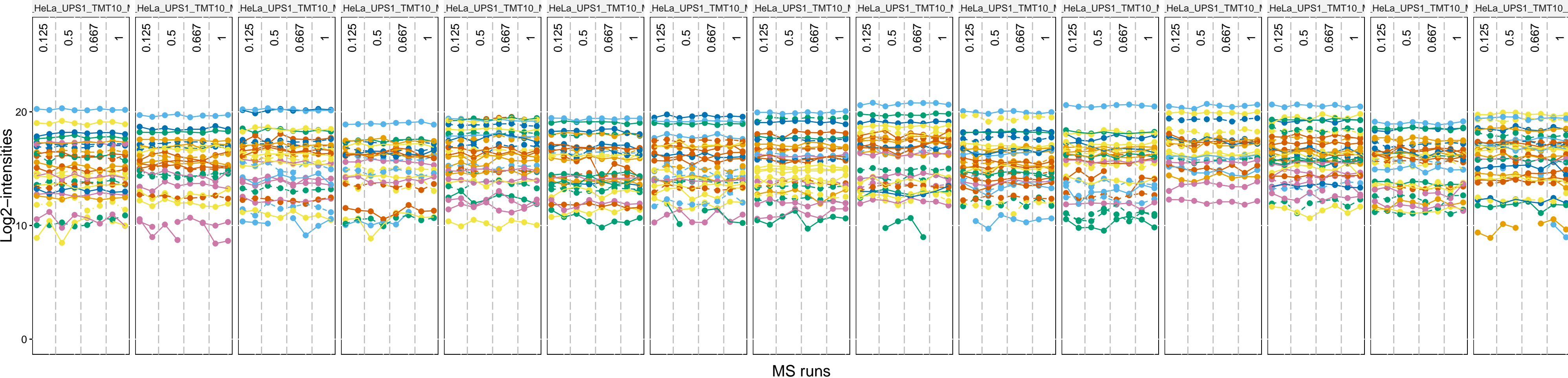
[R].trEGNDLYHEmIESGVINLk.[D]_4

[R].trEGNDLYHEmIESGVINLk.[D]_4

[R].vALTGLTVAEYFr.[D]_2

[R].vALTGLTVAEYFr.[D]_3

[R].vALTGLTVAEYFr.[D]_3



peptide: 19

- [K].dLFDPIEDr.[H]_2

[K].dYEFmWNPHLGyILTcPSNLGTGLr.[A]_3

[K].dYEFmWNPHLGyILTcPSNLGTGLr.[A]_4

[K].dYEFmWNPHLGyILTcPSNLGTGLr.[A]_4

[K].IAVEALSSLDGDLGr.[Y]_2

[K].IAVEALSSLDGDLGr.[Y]_3
- [K].ILIEMEqr.[L]_2

[K].rGTGGVDTAAVGGVFDVSNADr.[L]_3

[K].smTEAEQQQLIDDHFLFDk.[P]_3

[K].smTEAEQQQLIDDHFLFDk.[P]_3

[K].smTEAEQQQLIDDHFLFDk.[P]_4

[K].smTEAEQQQLIDDHFLFDkPVSPLLASGMAr.[D]_4
- [K].sTPSGFTLDDVIQTGVDNPGHPYImTVGcVAGDEESYEVfk.[D]_4

[K].sTPSGFTLDDVIQTGVDNPGHPYIMTVGcVAGDEESYEVfk.[D]_4

[K].sTPSGFTLDDVIQTGVDNPGHPYIMTVGcVAGDEESYEVfk.[D]_5

[K].tDLNPDNLQGGDDLDPNYVLSSr.[V]_3

[K].sMTAEAEQQQLIDDHFLFDk.[P]_3

[K].tFLVWVNEEDHLr.[V]_3

[K].vLTPELYAELr.[A]_2
- [R].fcTGLTQIETLfk.[S]_2

[R].fcTGLTQIETLfk.[S]_3

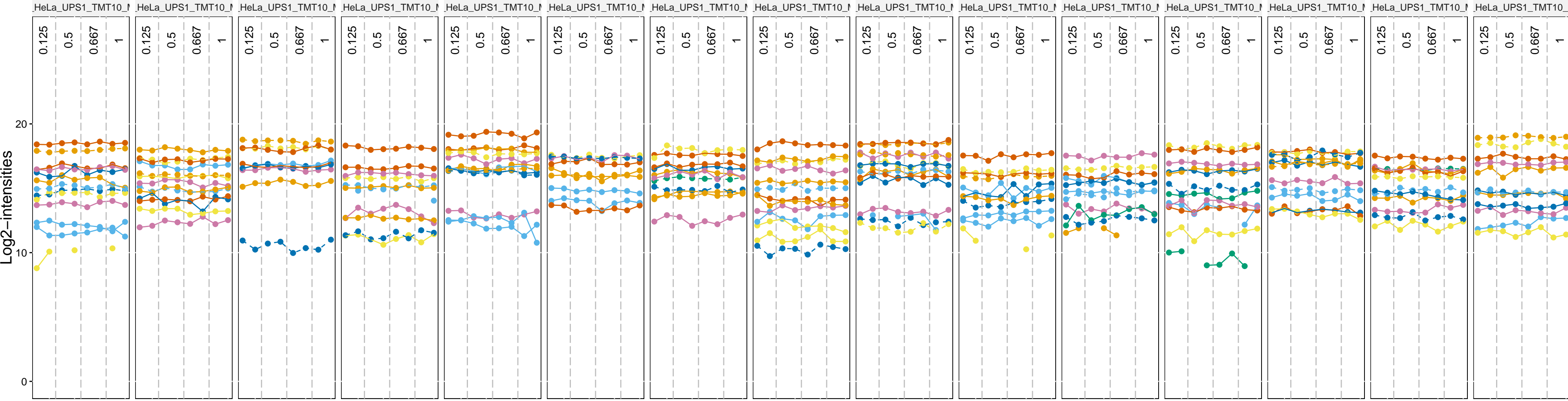
[R].gTGGVDTAAVGGVFDVSNADr.[L]_2

[R].gTGGVDTAAVGGVFDVSNADr.[L]_3

[R].IEQGQAIDDLmPAQk.[-]_3

[R].IGFSEVELVQMmVVDGVk.[L]_2
- [R].IGFSEVELVQmVVDGVk.[L]_3

[R].IGFSEVELVQMmVVDGVk.[L]_3



MS runs

peptide: 13

- [K].dTTLNWk.[M]_2

[K].eNFSLDWck.[Q]_2

[K].iLSSYLQk.[K]_2
- [K].IVEALcAAGHr.[A]_3
- [K].pLGELWk.[-]_2
- [K].qPDVGLPkPDLVLFLQLQLADAak.[R]_3

[K].qPDVGLPkPDLVLFLQLQLADAak.[R]_4

[K].sIEAVHEDlr.[V]_3

[R].aELLrFPEr.[S]_3

[R].gAFGHEr.[Y]_3

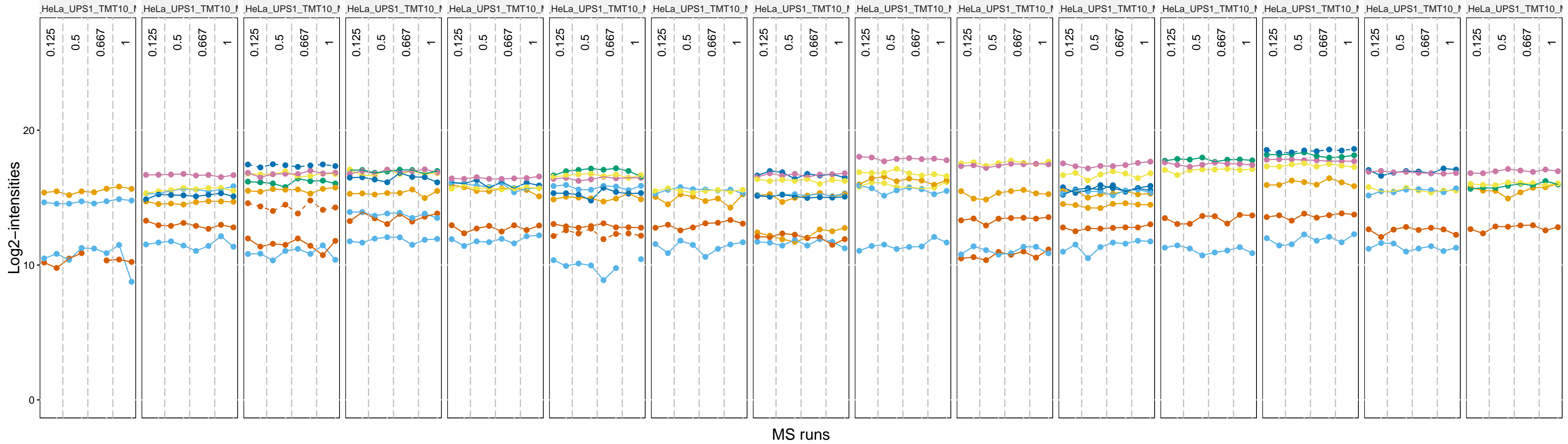
[R].rGALIVLEGVDr.[A]_3

[R].vLSEDAIr.[T]_2


















[R].yAFSGVAFTGak.[E]_2

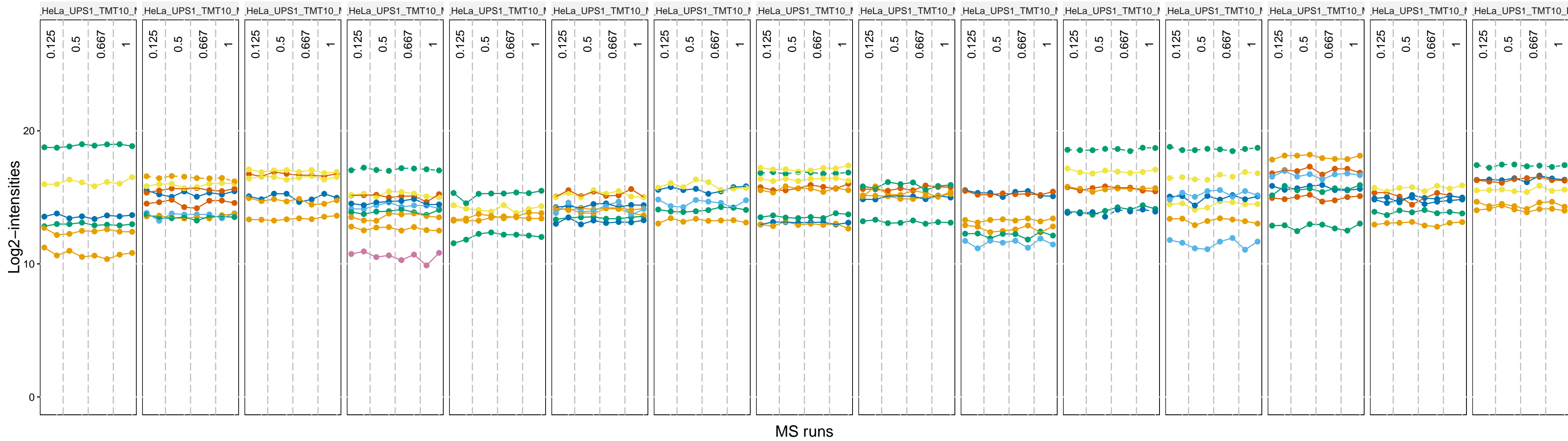
[R].yAFSGVAFTGak.[E]_3

[R].yENGAFQEr.[A]_2



peptide: 13

 [K].gEELScEEr.[N]_2	 [K].vETELQGVcDTVLGLLDShlik.[E]_3	 [R].sAYQEAMDISk.[K]_2	 [R].vLSSIEQk.[ST]_3	 [R].yLAEVATGDDkk.[R]_3
 [K].gEELScEEr.[N]_2	 [K].vETELQGVcDTVLGLLDShlik.[E]_4	 [R].sAYQEAMDISk.[K]_3	 [R].yEDMAAFMk.[G]_2	
 [K].rIIDSAr.[S]_3	 [R].dNLTlWTADNAGEEGEAPQEPQS.[-]_3	 [R].sAYQEAMDISkk.[E]_3	 [R].yEDMAAFMk.[G]_3	
 [K].tTFDEAMADLHLTSEDsYk.[D]_3	 [R].sAYQEAmDISk.[K]_2	 [R].vLSSIEQk.[ST]_2	 [R].yLAEVATGDDk.[K]_2	



Q15233

peptide: 36

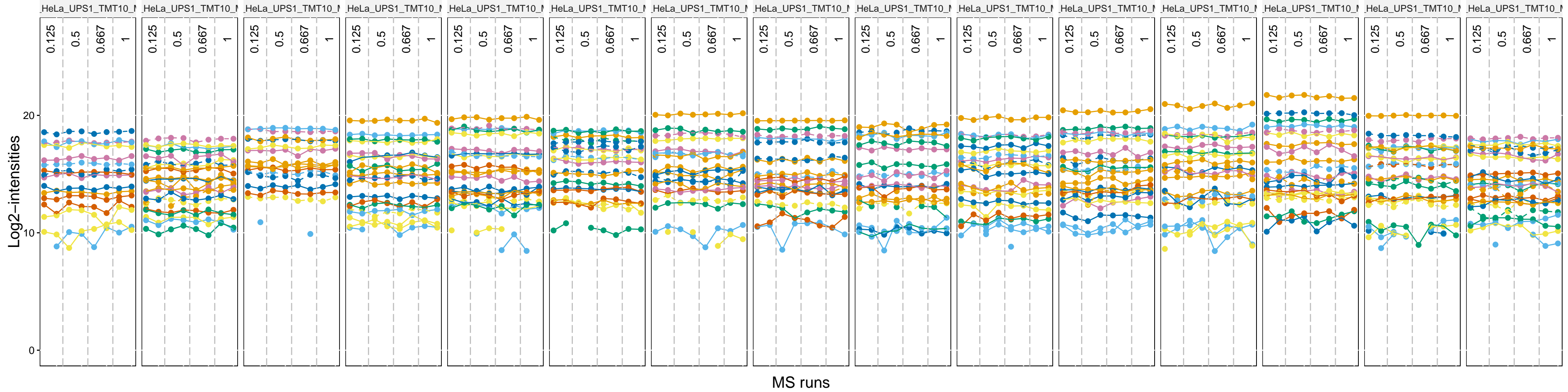
[K].aGEVFIHk.[D]_3
[K].aLIEMEk.[Q]_2
[K].gIVEFSGk.[P]_2
[K].gIVEFSGkPAAr.[K]_3
[K].iEmEMEAAr.[H]_2
[K].iEMEMEAAr.[H]_2
[K].iEMEMEAAr.[H]_2
[K].qLELrQEEEr.[R]_3
[K].qQQDQVDr.[N]_2
[K].vELDNmPLr.[G]_2

[K].vELDNMPLr.[G]_2
[R].aAPGAEFAPNk.[R]_2
[R].aAPGAEFAPNk.[R]_3
[R].cSEGSFLLTTFPr.[P]_3
[R].cSEGSFLLTTFPrPVTVEPmDQLDDEEGLPEk.[L]_4
[R].cSEGSFLLTTFPrPVTVEPmDQLDDEEGLPEk.[L]_4
[R].fAQPGSFYEYAmr.[W]_2
[R].fAQPGSFYEYAmr.[W]_2
[R].fAQPGSFYEYAmr.[W]_3
[R].fAQPGSFYEYAmr.[W]_3

[R].fGQAATMEGIGAIGGTPPAFNr.[A]_2
[R].fGQAATMEGIGAIGGTPPAFNr.[A]_3
[R].fGQAATMEGIGAIGGTPPAFNr.[A]_3
[R].gAMPPAPVPAGTPAPPGPATMMPDGTGLGTPPTTTr.[F]_3
[R].gAMPPAPVPAGTPAPPGPATmMPDGTGLGTPPTTTr.[F]_4
[R].gAMPPAPVPAGTPAPPGPATmMPDGTGLGTPPTTTr.[F]_4
[R].gAMPPAPVPAGTPAPPGPATMMPDGTGLGTPPTTTr.[F]_4
[R].hEHQVMLmr.[Q]_3
[R].hEHQVMLmr.[Q]_3

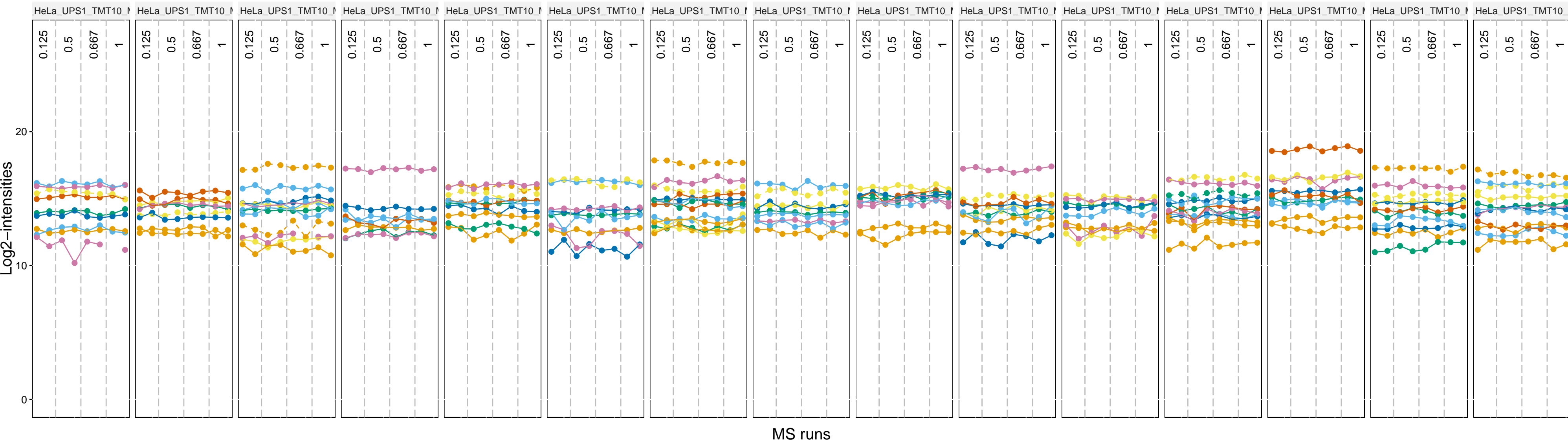
[R].hEHQVmLmr.[Q]_4
[R].hEHQVMLmr.[Q]_4
[R].hEHQVMLmr.[Q]_4
[R].hEHQVMLmr.[Q]_4
[R].iFVGNLPPDITEEEmr.[K]_2
[R].iFVGNLPPDITEEEmr.[K]_3
[R].iFVGNLPPDITEEEmrk.[L]_3
[R].mEELHNQEVQk.[R]_3
[R].mGQMAMGGAmGinnr.[G]_2
[R].mGQMAMGGAmGinnr.[G]_2

[R].mGQMAMGGAmGinnr.[G]_3
[R].nLPQYVSNELLEEAFSVFGQVER.[A]_3
[R].nLPQYVSNELLEEAFSVFGQVER.[A]_3
[R].qQEEMMr.[R]_2
[R].rMEELHNQEVQk.[R]_3
[R].rMEELHNQEVQk.[R]_4
[R].rQEEMMr.[R]_3



peptide: 16

	[K].aDTLTPEEcQQFk.[K]_2		[K].drLPLAVVGSNTIIEVNGk.[R]_3		[K].iYEFPETDDEEENk.[L]_2		[K].IAAVTYNGVDNNk.[N]_3		[K].sPLAQMEEErr.[E]_3
	[K].aDTLTPEEcQQFk.[K]_3		[K].dSEAEQLqr.[R]_2		[K].iYEFPETDDEEENk.[L]_3		[K].nLEGYVGfANLPNQVYr.[K]_2		[K].sTLINSLFLTDLYSPEYGPShr.[I]_3
	[K].aDTLTPEEcQQFk.[Q]_3		[K].dVTNNVHYENYr.[S]_3		[K].kMEMEMEQVFEMk.[V]_3		[K].nLEGYVGfANLPNQVYr.[K]_3		[R].iLEQQNSSr.[T]_2
	[K].aNWEAQqr.[I]_2		[K].fEDYLNAESr.[V]_2		[K].IAAVTYNGVDNNk.[N]_2		[K].sPLAQMEEErr.[E]_3		[R].qYPWGVAEVENGEHcDFTILr.[N]_3



Q9NSD9

peptide: 27

- [K].aAGASDVVLYk.[I]_2

[K].aSEGPAFFPGr.[C]_2

[K].drYDSFIELQEk.[L]_3

[K].eYTAcELMNIYk.[T]_2

[K].eYTAcELMNIYk.[T]_3

[K].hYLHIIENk.[P]_4

[K].iDVPANrYDLLcLEGLVr.[G]_3
- [K].irPFAVAAVLr.[N]_3
- [K].iFEISDIVIk.[D]_2
- [K].iFEISDIVIk.[D]_3
- [K].iGVDISATk.[A]_2
- [K].iGVLHPDVITk.[F]_3
- [K].iHQNIcr.[K]_3
- [K].IIITEETAk.[I]_2

[K].nPGFEIIHGLLDr.[I]_3

[K].rDLLFQALGr.[T]_3

[K].sHTFPelayr.[K]_3

[K].tAEFQVAr.[T]_2

[K].tYTIANQFPLNk.[L]_2

[R].aDIIHAcDIVEDAAIAYGYNNIQMTLPk.[T]_3

[R].aDIIHAcDIVEDAAIAYGYNNIQMTLPk.[T]_4

[R].aLVAIGTHDLDLTLGGPFTYTak.[R]_3

[R].aLVAIGTHDLDLTLGGPFTYTak.[R]_4

[R].eTPENLAK.[L]_2

[R].hDMAAAGFTTEALTfALcSQEDIADk.[L]_3

[R].hLcAVYYNk.[N]_3

[R].iMQLLDVPPGEDk.[G]_3

[R].kMPLPLk.[L]_3

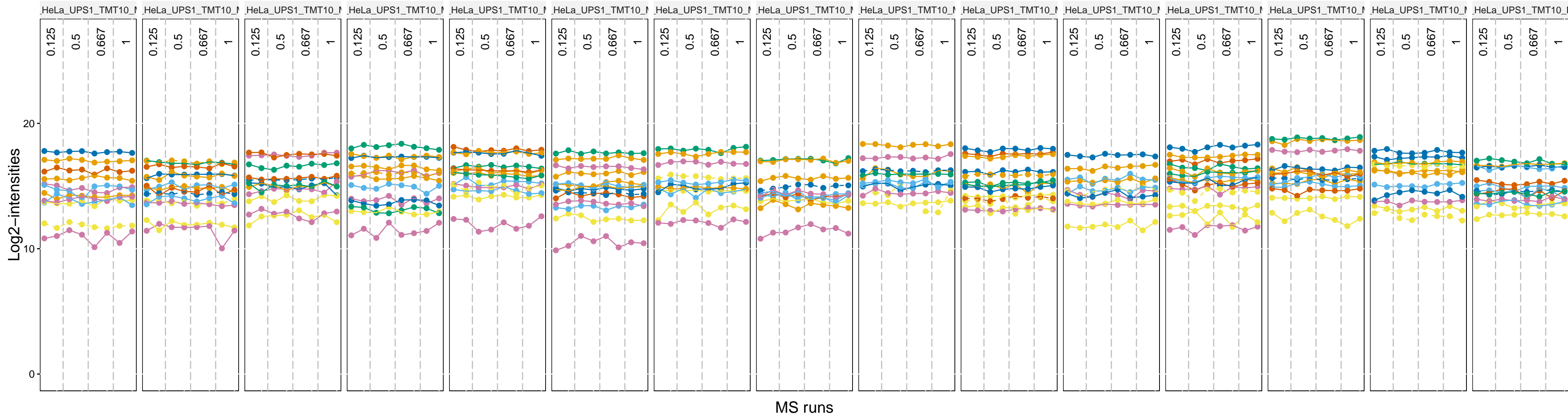
[R].nFIEcTGTDFtk.[A]_2

[R].tLLPGLLk.[T]_2

[R].tLLPGLLk.[T]_3

[R].yDLLcLEGLVr.[G]_2

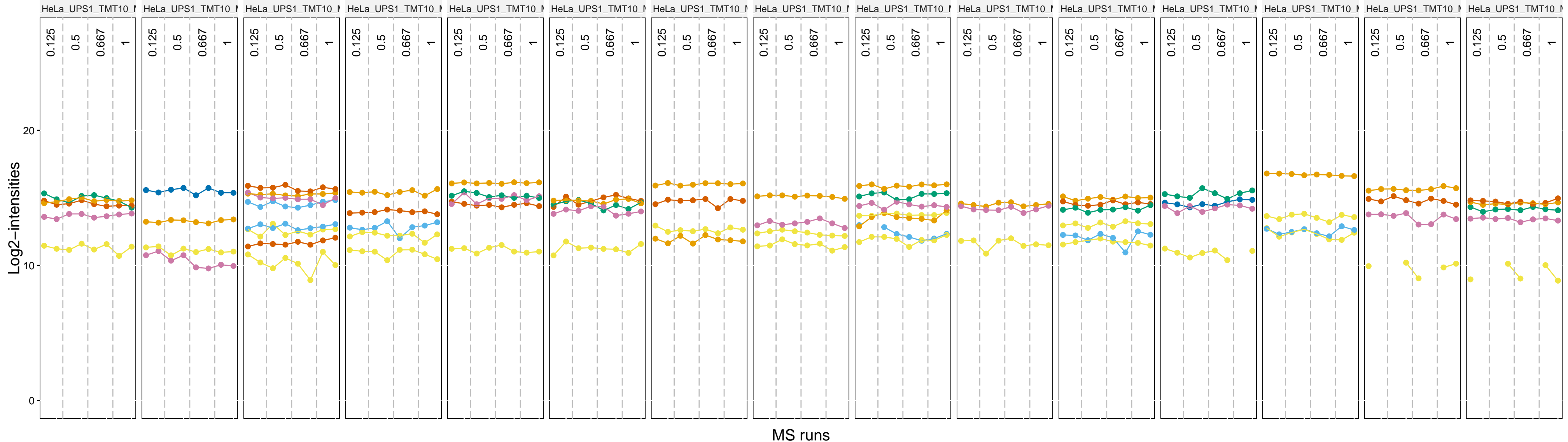
[R].yDLLcLEGLVr.[G]_3
















Q9UGP8

peptide: 15

- [K].aYAALTDEESr.[K]_2
- [K].eQSIcAAEEQPAEDGQGETNk.[N]_3
- [K].nWEEFGNPDGPQATSFGIALPAWIVDQk.[N]_4
- [K].tIQDLVSLk.[E]_2
- [R].eYQEYNPYEVLNLDPGATVAEIk.[K]_3
- [K].dATSrPTDNILIPQLr.[E]_3
- [K].gGDEVFMFr.[I]_2
- [K].qNkDDEAEWQELQQSIQr.[K]_3
- [R].dQNAEQIr.[L]_2
- [R].iIMVLAGASEFDPQYNk.[D]_3
- [K].dTEEVELk.[F]_2
- [K].kNEPPLTcPYSLk.[A]_3
- [K].sQVLDDSDNNITVGSLVTVLVk.[L]_3
- [R].eiGSINLk.[K]_2
- [R].vLLLSHLAr.[M]_3



peptide: 13

 [K].aLSGVLEQDr.[V]_2	 [K].gQNALVELQTQrPIALELYk.[D]_3	 [R].dkPSVEPVVEEYDYEDLk.[E]_3	 [R].gVTMDVGMTk.[F]_2	 [R].sESEIVPk.[V]_2
 [K].eSDVGFIPTSGLSGENLItr.[S]_2	 [K].iEAGYIQTDGr.[L]_2	 [R].eVLGDAVPDEILIEAVLk.[N]_3	 [R].iLIFNIEIPITk.[G]_3	
 [K].gPPIEDAIASSDVLETASK.[S]_3	 [K].rLISVLNk.[S]_3	 [R].gEFEAGFETGGQTr.[E]_2	 [R].lcVSDVFk.[D]_2	

