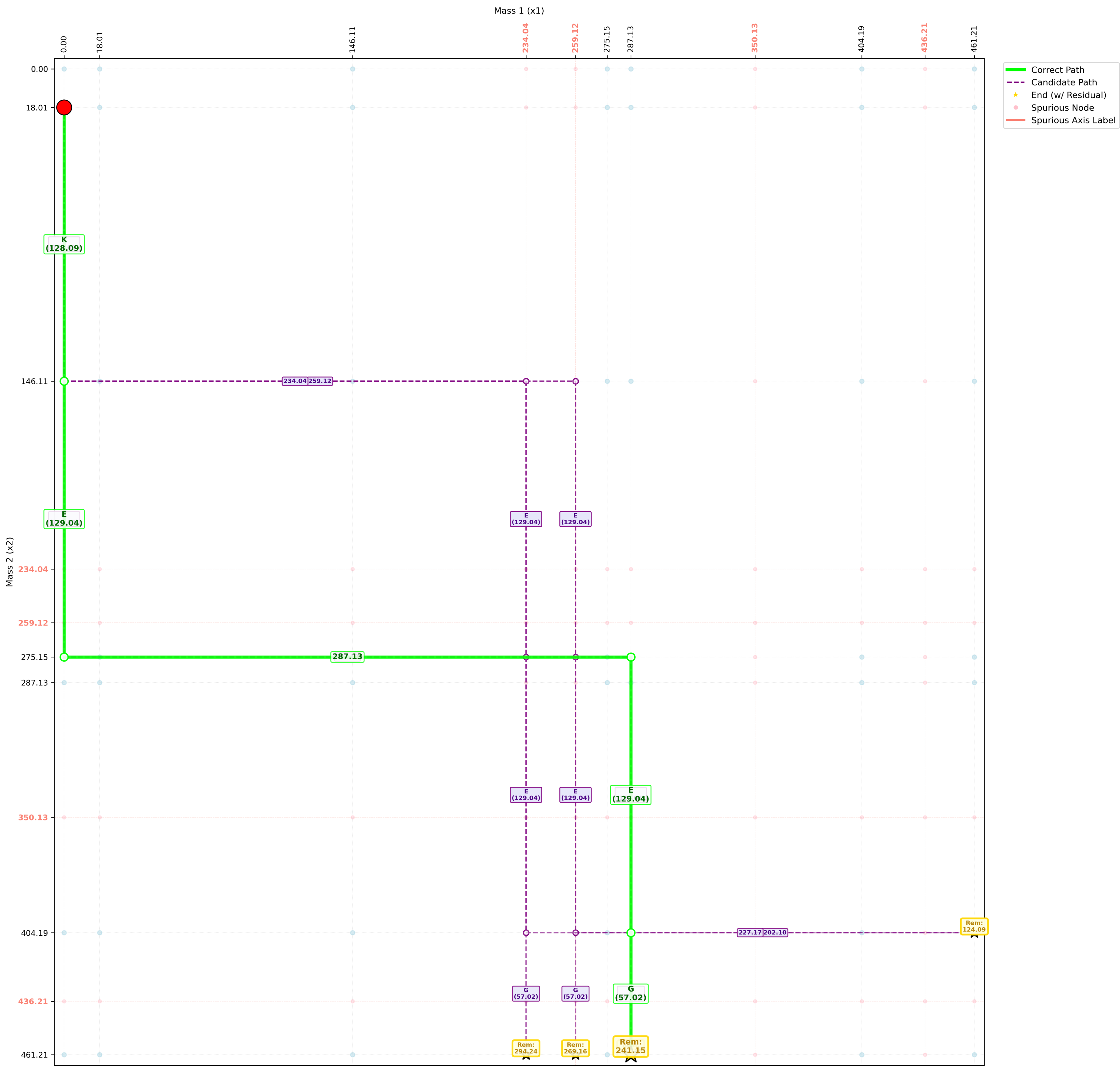


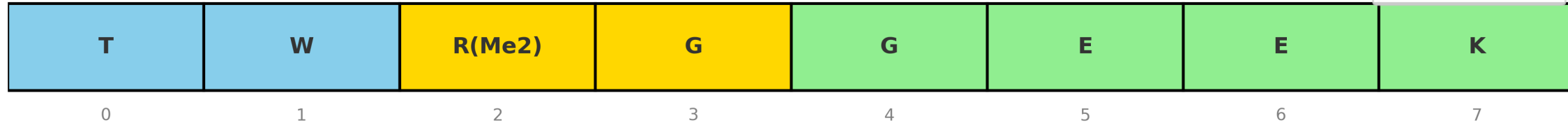
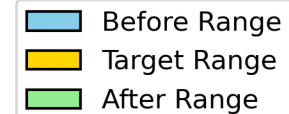
	<div> <div>b-ion</div> <div>y-ion</div> <div>Spurious</div> </div>																
	<div> <div>Peptide 1</div> <div>Peptide 2</div> <div>Peptide 3</div> <div>Peptide 4</div> <div>Peptide 5</div> </div>																
Ground Truth	146.105	234.041	259.116	275.148	287.127	350.131	404.191	436.207	461.212	471.196	471.259	518.234	528.281	585.302	714.345	843.387	971.482
	<div> <div>TW</div> <div>R(Me2)</div> <div>G</div> <div>G</div> <div>E</div> <div>E</div> <div>K</div> </div>																
Peptide 1			259.116						461.212					585.302	714.345	843.387	
Peptide 2			259.116										528.281	585.302	714.345	843.387	
Peptide 3		234.041											528.281	585.302	714.345	843.387	
Peptide 4		234.041							461.212					585.302	714.345	843.387	
Peptide 5					287.127								528.281	585.302	714.345	843.387	



ME17\_3+: [TWR(Me2)GGEEK+3H]3+



	b1y7(T)	b2y6(W)	b3y5(R(Me2))	b4y4(G)	b5y3(G)	b6y2(E)	b7y1(E)	Row_Count	Unexplained Count	Abs Average Mass Difference	Unexplained Pairs
Parent	--	--	(b3,y5) (2+ , 1+) (-0.18, 0.09) 1	(b4,y4) (2+ , 1+) (-0.1, -0.02) 5	(b5,y3) (2+ , 1+) (-0.17, -0.22) 31	(b6,y2) (2+ , 1+) (-0.16, 0.04) 3	(b7,y1) (2+ , 1+) (-0.43, 0.11) 25	5	1	0.3	-0.43(28)
Col_Count	0	1	3	2	2	2	3	13	1	0.14	nan
Ion Mass	(102.05, 889.45)	(288.13, 703.37)	(472.27, 519.24)	(529.29, 462.22)	(586.31, 405.2)	(715.35, 276.16)	(844.39, 147.11)	0	0	0.0	0

Array Visualization (Target Range: [2 - 3])



 b-ion     y-ion     Middle Point

146.105

275.148

287.127

404.191

461.212

287.127

461.212