

Full Scan	
Resolution	70,000 (@ m/z 200)
AGC target value	3e6
Scan range	350-2000 m/z
Maximal injection time (ms)	100
Data-dependent MS/MS	
Inclusion	Off
Resolution	17,500 (@ m/z 200)
AGC target value	1e5
Maximal injection time (ms)	50
Loop Count	10
Isolation window width (m/z)	2
NCE (%)	27
Data-dependent Settings	
Underfill ratio (%)	1
Charge exclusion	Unassigned, 1, 7, 8, >8
Peptide match	preferred
Exclusion isotopes	on
Dynamic exclusion (s)	60