

Parameter	Value
Number of combined spectra	1
Fragmentation method	FILE
Search type	TARGET+DECOY
Fixed modifications	None
Maximum number of unexpected modifications	0
Error tolerance	15 ppm
Cutoff type	FDR
Cutoff value	0.01
Allowed N-terminal forms	NONE,NME,NME_ACETYLTATION
Maximum mass shift of modifications	500 Da
Thread number	1
E-value computation	Lookup table