

Correlation settings	
Insertion of new peptides <ul style="list-style-type: none"> do not insert new peptides insert peptides only in range 	checked unchecked
LC peak detection <ul style="list-style-type: none"> left LC window (secs) right LC window (secs) 	30 30
„use fixed“	Wild type
Options	
Include in preselected peptides (PSPs) <ul style="list-style-type: none"> bold red light red, bold black parenthesized peptide filter 	checked checked checked checked Mascot score 1 -200
Recalibration of preselected peptides <ul style="list-style-type: none"> domain peptide filter 	Frequency, 1/ ^{0.5} (OrbiTrap) Mascot score 30 - 200
Validation of preselected peptides <ul style="list-style-type: none"> peptide filter 	Mascot score 15 - 200
Classification of preselected peptides (PSPs) by Mascot score <ul style="list-style-type: none"> category A category B category C protein is validated by default if ABC score is greater than or equal to 	15 15 15 30
Quantitation mode	v2.2 ICAT 9C
MGF file generator	NewDTASupercharge, up to 1.18....
Raw file type	LTQ-FT/LTQ OrbiTrap (.raw)
Type(s) for quantitation	all MS spectra

SUPPLEMENTAL TABLE 1. MSQuant parameters used for quantification of LC-MS/MS data