$S$ comments ← All the lines starting with "$S" are headers.
FUNCTION 1← Indicates the beginning of MS-1 raw data.
Scan 1← Indicates the beginning of the precursor scan and its serial number.
Retention Time 0.026← Indicates the retention time of the LC and the beginning of the MS data.
449.5780 0← MS raw data (m/z value and intensity value).
449.7510 0
:
Scan 2← The next scan and its serial number.
:
FUNCTION 2← Indicates the beginning of MS-2 raw data (the first data acquisition).
Scan 1
Retention Time 25.479← Indicates the retention time of the LC and the beginning of the MS/MS data.
Set Mass 459.2822← The peak m/z value of the selected parent peptide.
Collision Energy 30.0000
49.5864 0← MS/MS raw data (m/z value and intensity value).
70.0673 0
70.0707 1
:
FUNCTION 3← Indicates the beginning of MS-2 raw data (the second data acquisition).

: