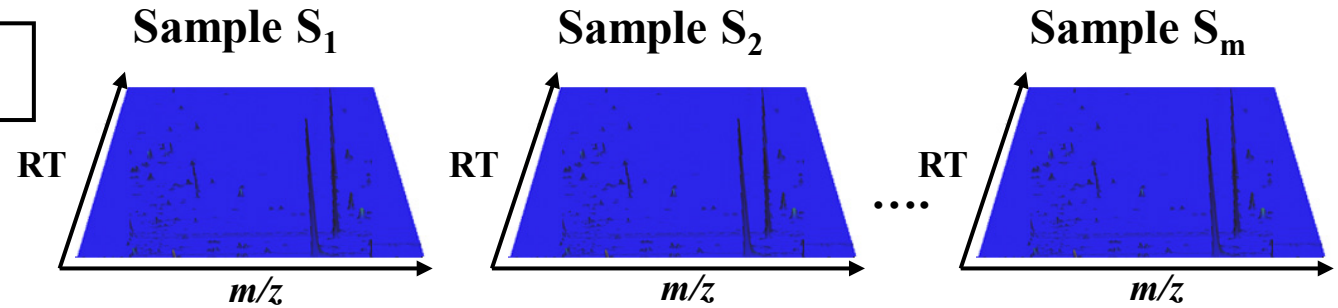
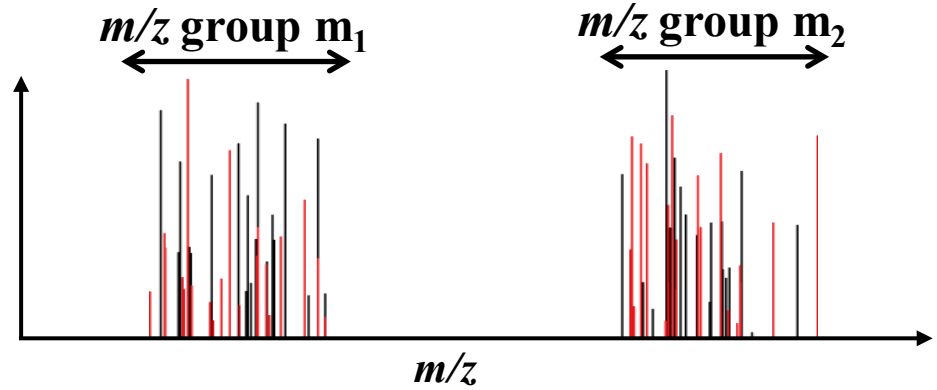
(a) Collect data points



(b) Group collected data by *m/z* closeness



(c) Extract ion chromatograms for *m/z* list

m/z	S_1	S_2			S_m
m_1	x_{11}	x_{12}			x_{1m}
m_2	x_{21}	x_{22}			x_{2m}
m_n	x_{n1}	x_{n2}			x_{nm}