

PepPre: Promote Peptide Identification



& using accurate and comprehensive precursors & bridge between DIA data and DDA engines



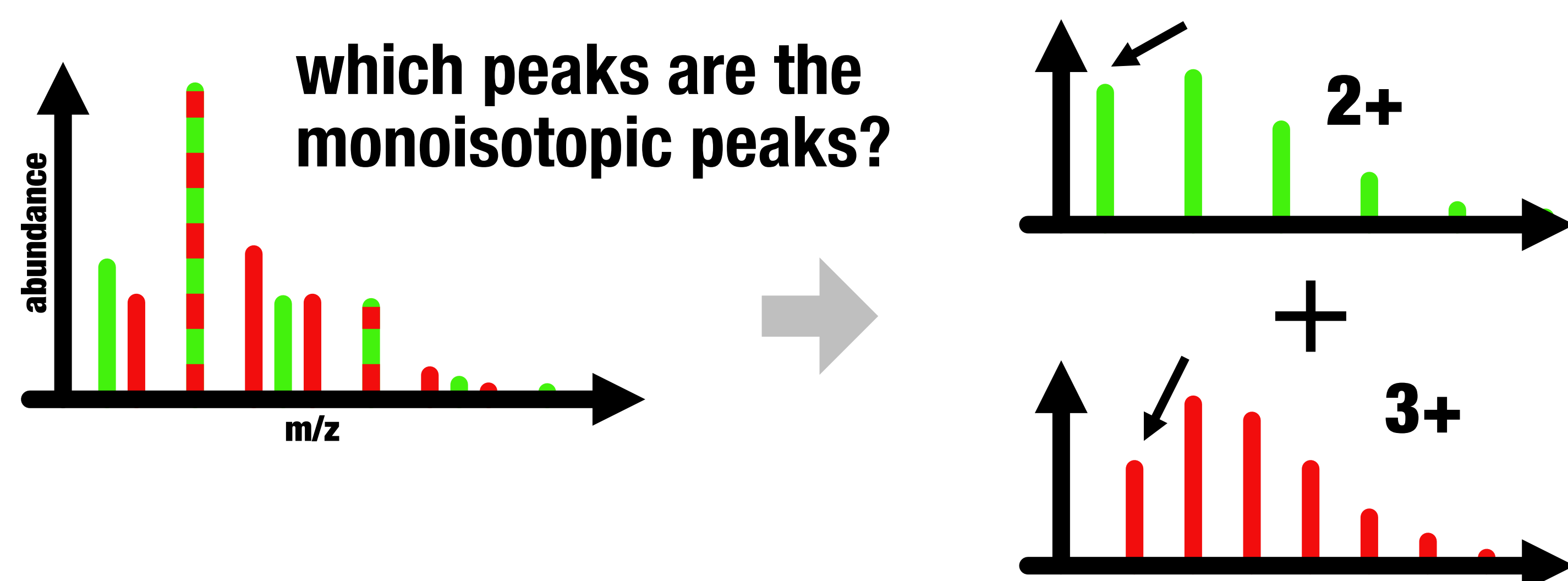
Introduction

Accurate and comprehensive **peptide precursor ions** are crucial to tandem mass spectrometry-based peptide identification.

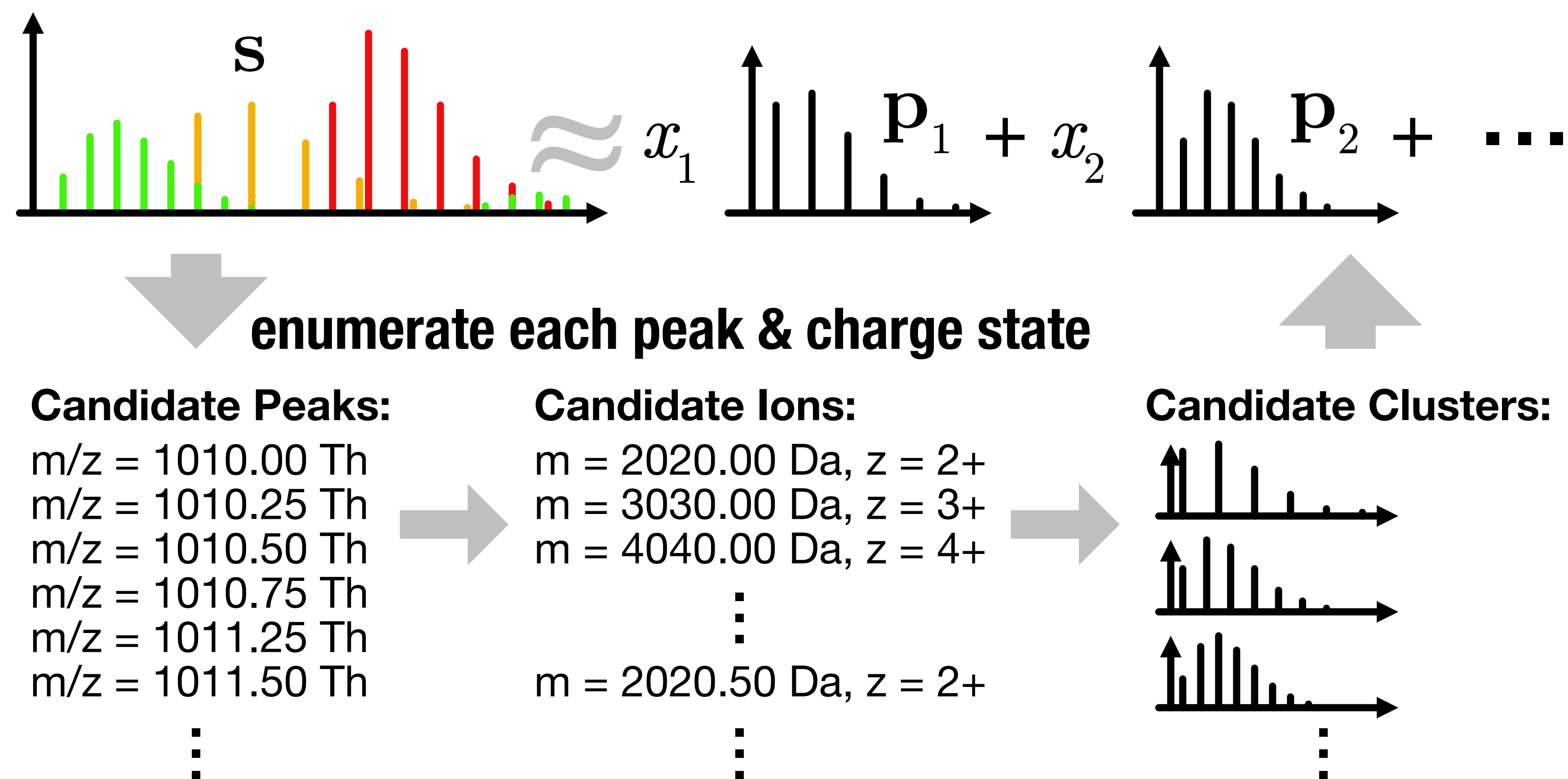
- An identification engine can derive great advantages from the **search space reduction** hinted by credible and detailed precursors.
- By considering multiple precursors per **mixed spectrum**, both the number of identifications and the spectrum explainability can be substantially improved.

Methods

Question: accurate mass & charge state of a peptide



Model: a spectrum is a vector



Solution: linear programming

$$s \approx x_1 p_1 + x_2 p_2 + \dots + x_n p_n \quad (1)$$

$$\Delta = |s - \sum_{i=1}^n x_i p_i| \quad (2)$$

$$\begin{aligned} \min \quad & \Delta \\ \text{s.t.} \quad & |s - \sum_{i=1}^n x_i p_i| = \Delta \\ & x_i \geq 0 \quad i = 1, 2, \dots, n \end{aligned} \quad (3)$$

Ching Tarn / 谭永卿, Email: i@ctarn.io, WeChat: TarnYeongChing
 pFind Team, Institute of Computing Technology, Chinese Academy of Sciences / 中国科学院计算技术研究所pFind课题组
 Made for 12th CNHUPO, 2023.9.24-27, Chengdu / 成都, China / 中国
 Tarn, C. et al. (2023). PepPre: Promote Peptide Identification Using Accurate and Comprehensive Precursors. bioRxiv, 2023.2005.2013.540645. <https://doi.org/10.1101/2023.05.13.540645>

Applications

Promote DDA Peptide Identification

HeLa, 2 hours, no fraction

Normal Isolation Window (± 1 Th):

Baseline:

#PSM = 39794

#Pep. = 28635

PepPre:

#PSM = 72934 (+83%)

#Pep. = 45341 (+58%)

Wide Window (± 4 Th):

Baseline:

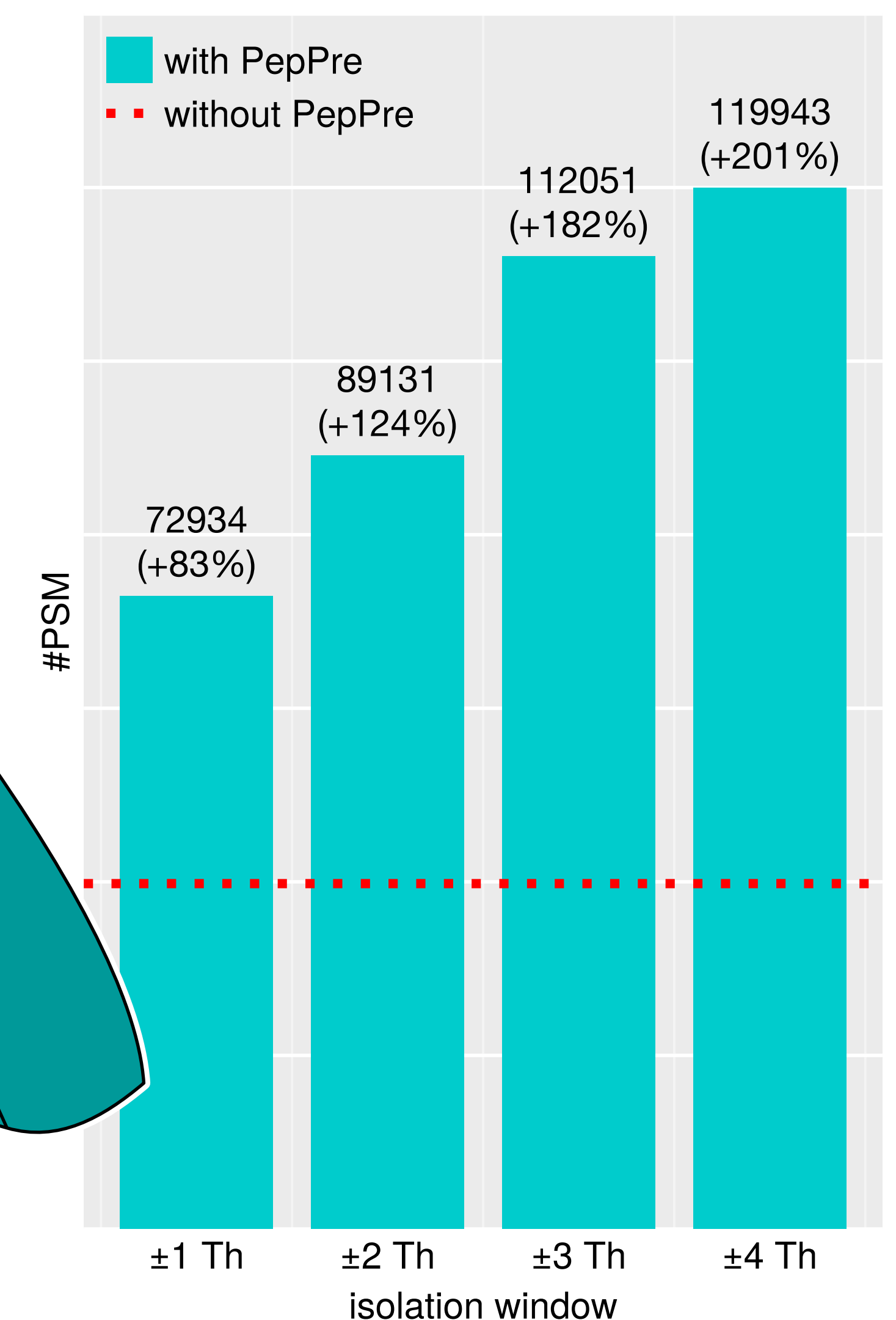
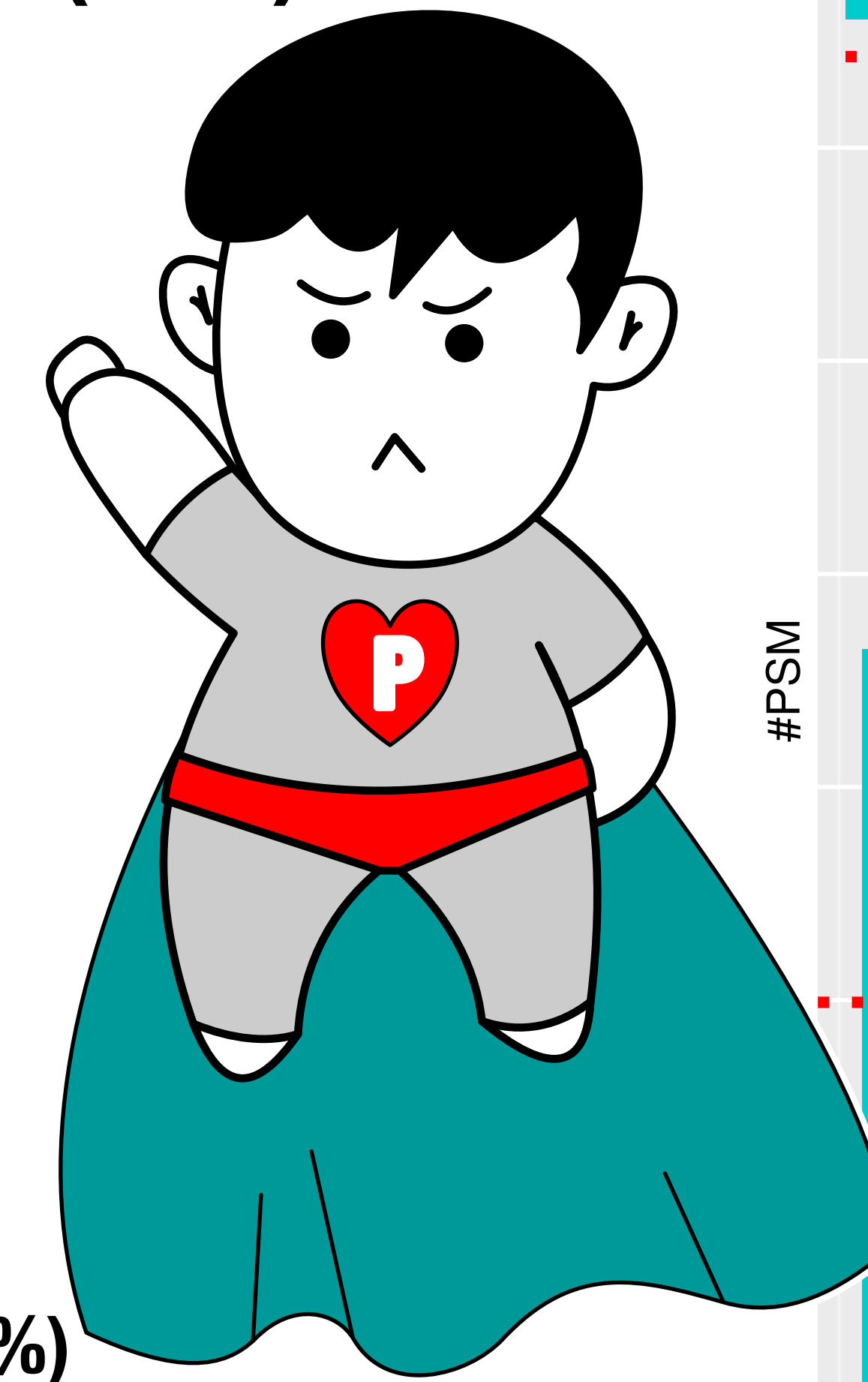
#PSM = 39590

#Pep. = 28255

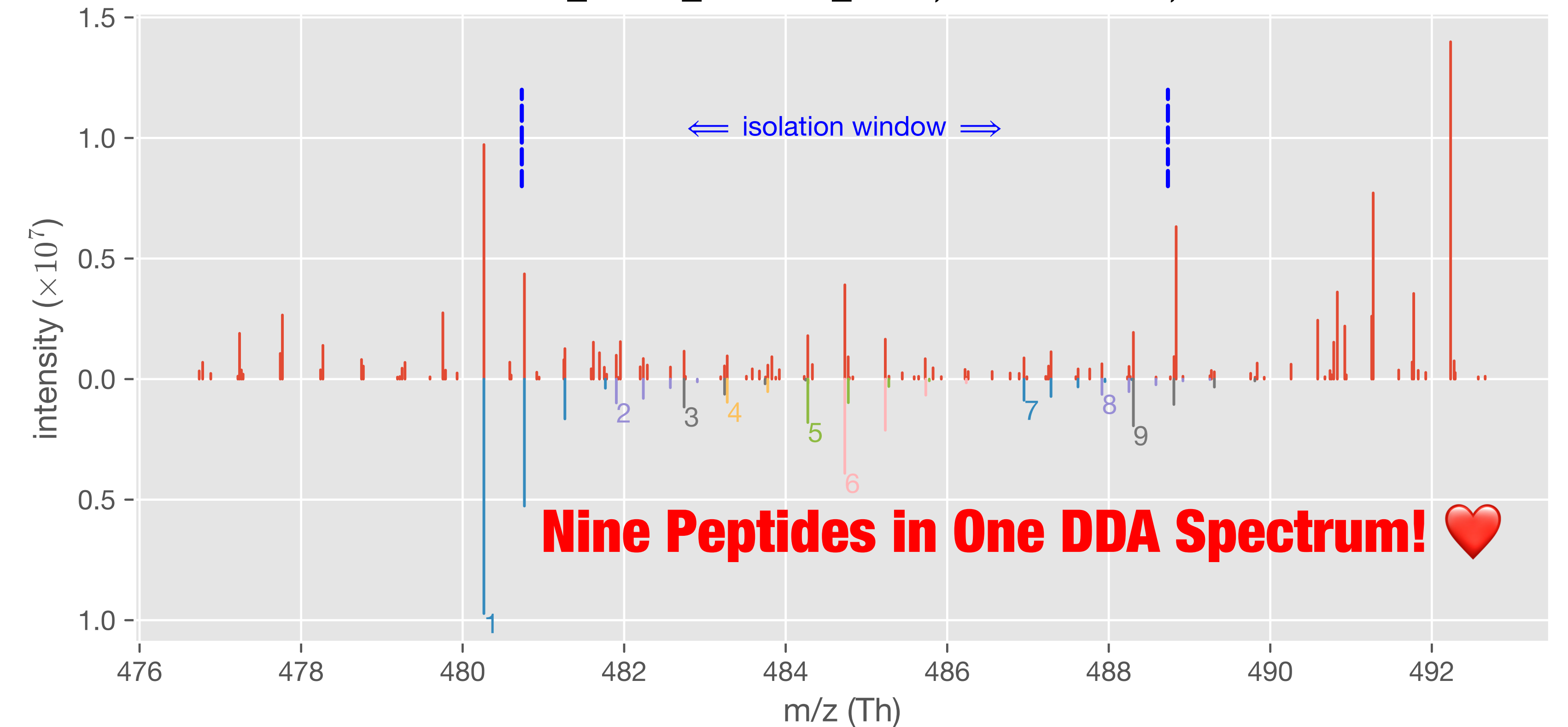
PepPre:

#PSM = 119943 (+203%)

#Pep. = 47370 (+65%)



RAW: 20131202_hela1_120min_8MZ, MS1: 19937, MS2: 19942



	m/z (Th)	charge state	sequence	modification	q-value
#1 ion	480.263305	2	SLEVLNLR	-	0.08%
#2 ion	481.903038	3	SEPHSLSSEALMR	-	0.11%
#3 ion	482.742083	2	FEELAAAR	-	0.36%
#4 ion	483.276243	2	LLSQVSYR	-	0.20%
#5 ion	484.273943	2	KEFFLQR	-	0.46%
#6 ion	484.731832	2	NGDYLCVK	Carbamidomethyl(C)@6	0.02%
#7 ion	486.950699	3	KPLPDHVSLEPK	-	0.14%
#8 ion	487.915765	3	LSNRPAFMPSEGR	-	0.45%
#9 ion	488.305021	2	KGVALNFVK	-	0.11%

Bridge between DIA data and DDA engines



<http://peppre.ctarn.io>

