

K9me3K14ac,+2, 15.2, 2.7e+06

20

26

Time (min)

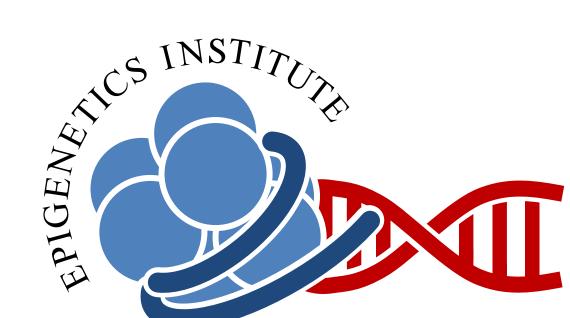
Figure 4. H3 9–17 KSTGGKAPR in endogenous samples.

30

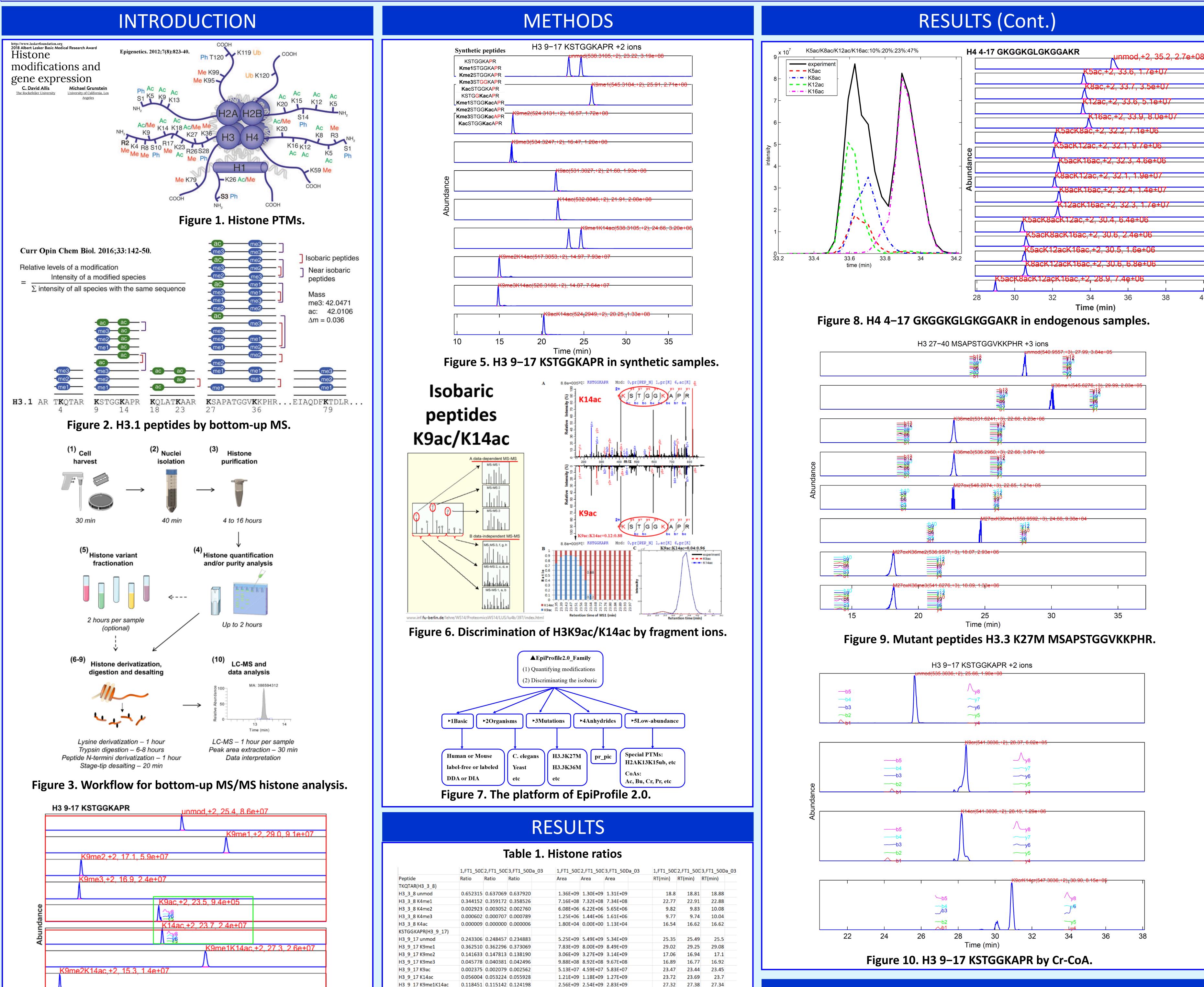
32

EpiProfile 2.0: A Computational Platform for Processing Epi-Proteomics Mass Spectrometry Data

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5.14E+08 5.34E+08 5.21E+08

9.19E+07 1.01E+08 9.69E+07

4.66E+10 4.79E+10 4.81E+10

9.40E+07 1.01E+08 1.08E+08

8.69E+07 8.90E+07 9.38E+07

5.61E+07 5.18E+07 5.98E+07

3.35E+09 3.31E+09 3.28E+09

1.76E+10 1.81E+10 1.81E+10

2.41E+09 2.33E+09 2.23E+09

15.18 15.33 15.26

39.07 39.15 38.98

41.47 41.47 41.42

42.22 42.29 42.28

43.3 43.34 43.36

37.23 37.31 37.1

37.38 37.5 37.25

35.34 35.5 35.21

H3_9_17 K9me2K14ac

H3_9_17 K9me3K14ac

KQLATKAAR(H3_18_26)

H3_18_26 unmod

H3_18_26 K23me1

H3_18_26 K18me1

H3_18_26 K18ac

H3_18_26 K23ac

0.004257 0.004594 0.004261

0.663978 0.666229 0.668268

0.001339 0.001409 0.001503

0.001238 0.001236 0.001303

0.047779 0.046073 0.045651

0.250533 0.251979 0.251394

H3_18_26 K18me1K23me1 0.000799 0.000720 0.000831

H3 18 26 K18acK23ac 0.034333 0.032353 0.031051

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