Peptide Mapping Sample to knowledge easier and faster than before

Peptide maps require 100% sequence coverage to prove the molecular structure of the biotherapeutic



Faster

Therm

Using the Thermo Scientific[™] SMART Digest[™] kit

thyroglobulin is digested in 3.5 hours versus 20 hours with in-solution digestion

Typical Digestion Times

Protein	Digest Time (min)
Insulin	4
BSA	< 5
Carbonic anhydrase	< 5
Lysozyme	< 5
Аро-В	30
lgG	45
lgG in 50 µL plasma*	75



Ribonuclease A 150 240 Thyroglobulin

C-reactive protein

*lgG in plasma (17.5 mg/mL total protein) at 70 °C

240

Reproducibility

The **SMART Digest kit** is highly reproducible user-to-user. Peak area RSD of...



Overlay of 3 separate digestions of a mAb using the SMART Digest kit (15mins), by 3 individual users



Automation

Las Automate Samples

Digestion of samples can be easily automated with the Thermo Scientific[™] KingFisher[™] purification systems, allowing for fast, high throughput sample processing. KingFisher systems help you to reduce hands on time and bring reproducibility to your workflow.

Separation

Highly reproducible



Peptide mixture: **Sequence:** Formula: Exact mass:

[Val5]-Angiotensin II **DRVYVHPF** $C_{49}H_{69}N_{13}O_{12}$ $[M+2H]^{2+} = 516.76671$

516.65 516.80 516.85 516.70 516.75 516.90 Mass (amu) [Lys-des-Arg⁹]-Bradykinin **KRPPGFSPF** $C_{50}H_{73}N_{13}O_{11}$ $[M+2H]^{2+} = 516.78490$ Joshua J. Coon, et al. ASMS 2012 oral, MOB pm

Dm (mmu):

18.2 mmu



Data Interpretation

Easy data interpretation



200.9% MS2/Fuil 17.71 712.985 200.9% MS2/Fuil 38.28 705.982 200.9% MS2/Fuil 38.28 1054.983

Thermo Scientific[™] BioPharma Finder[™] software allows easy interrogation of peptide maps enabling full characterization and confidence in results

Rituximab light chain

Rituximab heavy chain



For more information: thermofisher.com/SMARTdigest For more information on peptide mapping: thermofisher.com/peptidemaps For technical information: thermofisher.com/chromexpert

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