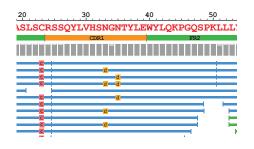
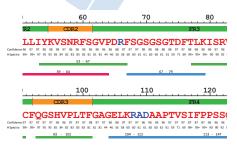
Reporting

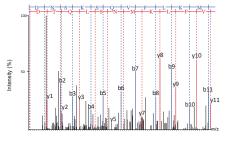
Once our team has completed and verified the antibody sequence, we provide a detailed, comprehensive report, which includes key information, such as:

- Constructed sequences and extensive peptide mapping
- Raw experimental LC-MS/MS data
- Sequence-level validation using Intact Mass Measurement
- Protein sequence confidence assessment
- Annotated supporting spectra
- Leucine and Isoleucine Differentiation
- And much more!



Leu/Ile Statisical Table				
Position	% of I	% of L	# of Matches	
1@3	36%	0%	0	
L@9	0%	96%	27	
L@11	0%	100%	10	
L@15	8%	91%	13	
I@21	100%	0%	3	
1@29	43%	56%	0	
L@38	1%	96%	20	





The best part is? Our turnaround time is less than 2 weeks. Ask us about our First-Time Customer discount and Batch-Order Pricing!

BIOINFORMATICS SOLUTIONS, INC.

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Tel: (519) 885-8288 Fax: (519) 885-9075

peaksab@bioinfor.com www.bioinfor.com





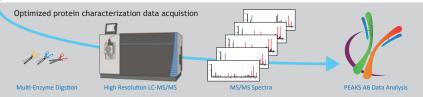


EXPERTISE YOU CAN TRUST SERVICE YOU CAN RELY ON



PEAKS AB Service delivers accurate *de novo* sequencing of monoclonal antibody (mAb) proteins. Aside from the correct protein sequence, PEAKS AB also examines intact mass measurement for sequence validation, and quantification of post-translational modifications (PTMs)/sequence variants upon request. Our PEAKS AB Service guarantees 100% sequence coverage and 100% accuracy of the constructed protein sequences of heavy and light chains.

Antibody Sequencing Using LC-MS/MS



Protein de novo sequencing powered by PEAKS AB Software &

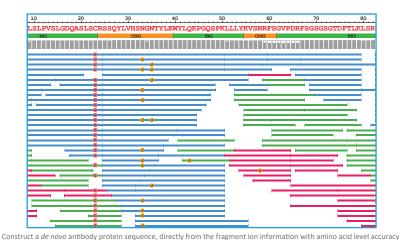
 DVLMTIQTPUSLIG
 DVLEVI LQLPHEBLEPUSLIG

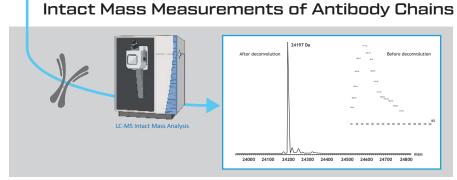
 STEPVISLIGNOJASILSEÇRISSEQY
 STEPVISLIGNOJASILSEÇRISSEQY

 SSQYLLVHISNIGNTYLIEWYLLQIKPGQSEPK
 STEPVISLIGNOJASILSEÇRISSEQY

 QKPGQISPKLLLLY
 QKPGQISPKLLLY

Peptide de novo sequencing Automated sequence assembly



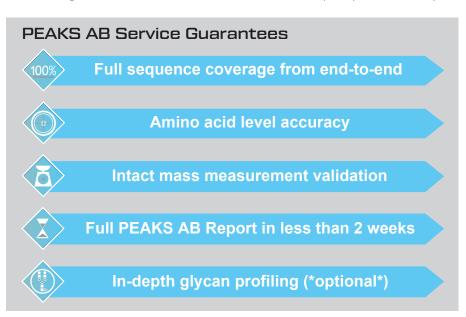


Assembled antibody sequence is validated with intact measurement of light chain and heavy chain by LC-MS

PEAKS AB Monoclonal Antibody Sequencing Details

Time	You Provide	We Deliver
2 weeks	Monoclonal antibody sample (>0.1mg, >80% purity)	 Full protein sequences 3-tier Leu/lle Differentiation Comprehensive PEAKS AB report Raw LC-MS/MS data Interactive result viewer

Price match guarantee and discount available for academic institutions or multiple samples. Contact us today!



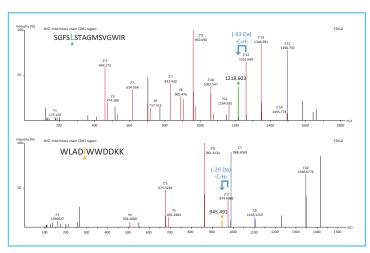
Confident Leucine & Isoleucine Differentiation

Overall performance of the antibody's specificity and affinity can be assured with Confident Leucine and Isoleucine Differentiation.

In our Antibody Sequencing Service, we use an integrated 3-tier strategy that combines:



- ① w-ion detection in
- enzyme cleavage preference.
- ③ homology statistics for unambiguous discrimination of Leu/Ile residues.



Characteristical loss of C_3H_7 (-43 Da) or C_2H_5 (-29Da) from the side chain of particular z-ion allows the Leu/Ile discrimintation