

What we can do for you:

- Peptides <30 aa routine; ~60-70 aa on request
- Amounts: typically <10 mg; higher amounts (<500 mg) feasible on request
- Solid phase Fmoc methodology
- HPLC purification if requested/required
- MS (ES, MALDI-TOF) validation of structure
- Conjugation to carrier protein, affinity columns, etc.

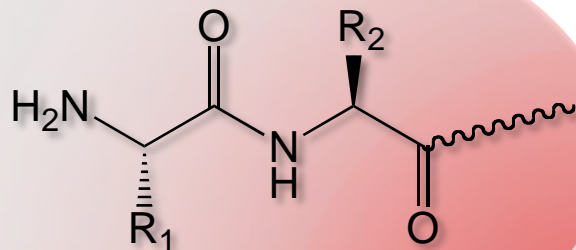
Typical workflow for an average peptide sequence



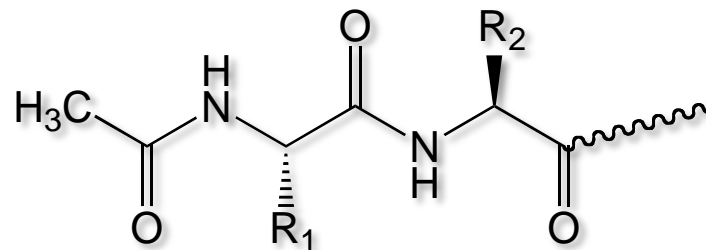
•2009-2014: ~125 jobs/yr (of which ~60% PRBB; external users include Spanish and EU public labs + small pharma & biotech companies)

End group options

N-terminus

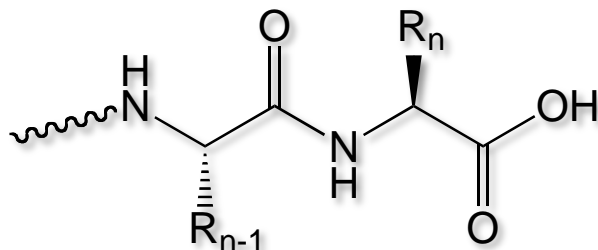


Free ($^+\text{NH}_3^-$ in solution)
Usually good for solubility
Required if actually native N-terminus

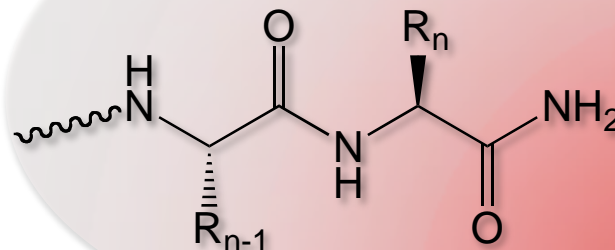


Acetylated (no charge)
Mimics better native situations

C-terminus



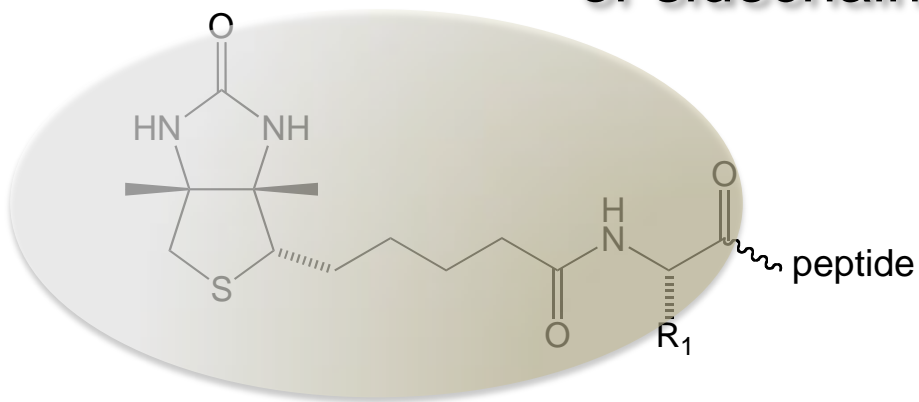
Free ($-\text{COO}^-$ in solution)
Required if actually native C-terminus



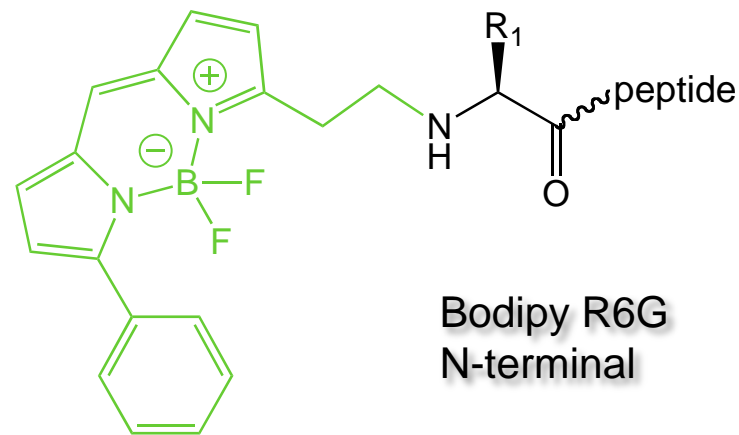
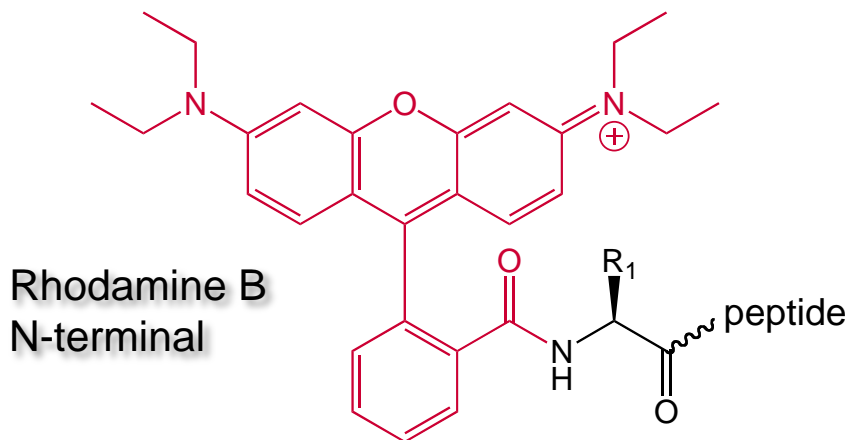
Carboxamide (no charge)
Mimics better native situation
Synthetically preferred

Site-specific modifications (1)

Biotinylation (N-terminal, C-terminal
or sidechain (Lys))

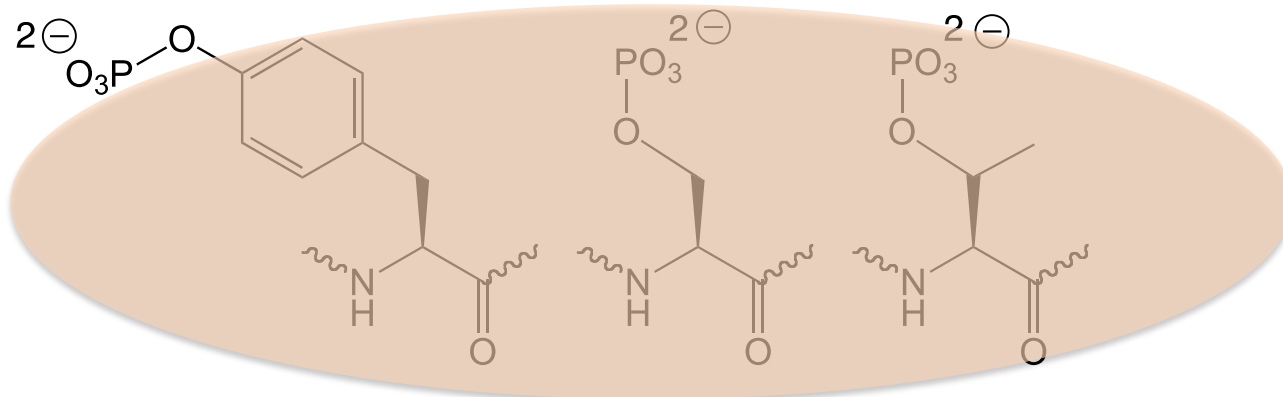


Fluoro labeling (different dyes)

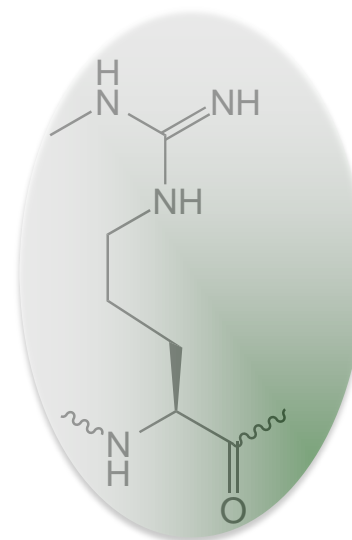
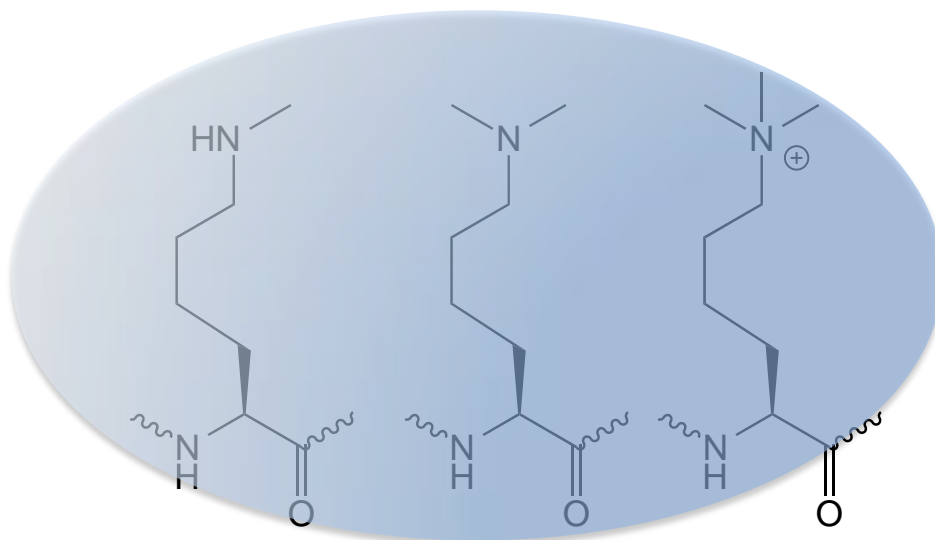


Site-specific modifications (2)

Phosphorylation (Tyr, Ser, Thr), sulfation (Tyr)



Modified Lys (methyl, acetyl) & Arg \rightarrow epigenetic studies



Other modifications, on request (3)

- D-Amino acids
- Non-coded amino acids
- Isotopically labeled (non-radioactive) amino acids (^{13}C , ^2H , etc.)
- Lipidation (fatty acids, farnesyl) at selected (Cys, Lys) positions
- Internal disulfides
- Complex, folded (multiple SS) peptides, protein mini-domains

Facility equipment

A. Synthesizers



PTI Prelude
Current workhorse



CEM Liberty Blue
In service, July 2015



ABI 433A (2001-2013)
phased out

Facility equipment (2)

B. HPLC*



preparative purification

C. HPLC-MS*



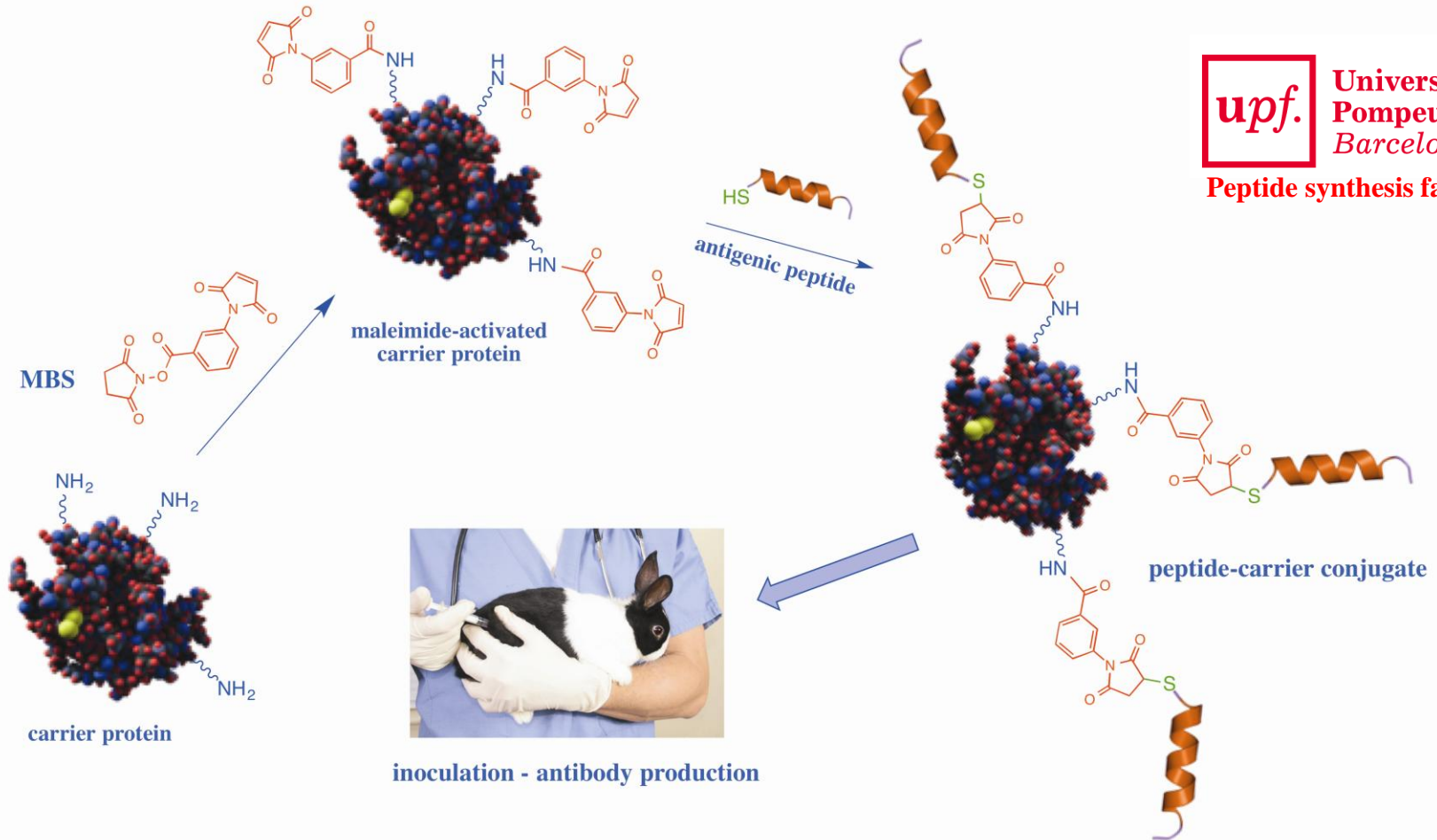
structural validation



analytical QC (3 systems)

*equipment available from the UPF proteomics and protein chemistry laboratory

Conjugation to carrier proteins or affinity supports



- An additional Cys residue added N-terminal or C-terminal to enable conjugation
- KLH as typical carrier (also BSA, OVA, TT)
- Heterobifunctional (maleimide-based) coupling chemistries ensures efficiency/selectivity
- AAA of conjugate to determine peptide/carrier molar ratio
- Similar chemistry for peptide-displaying affinity columns