

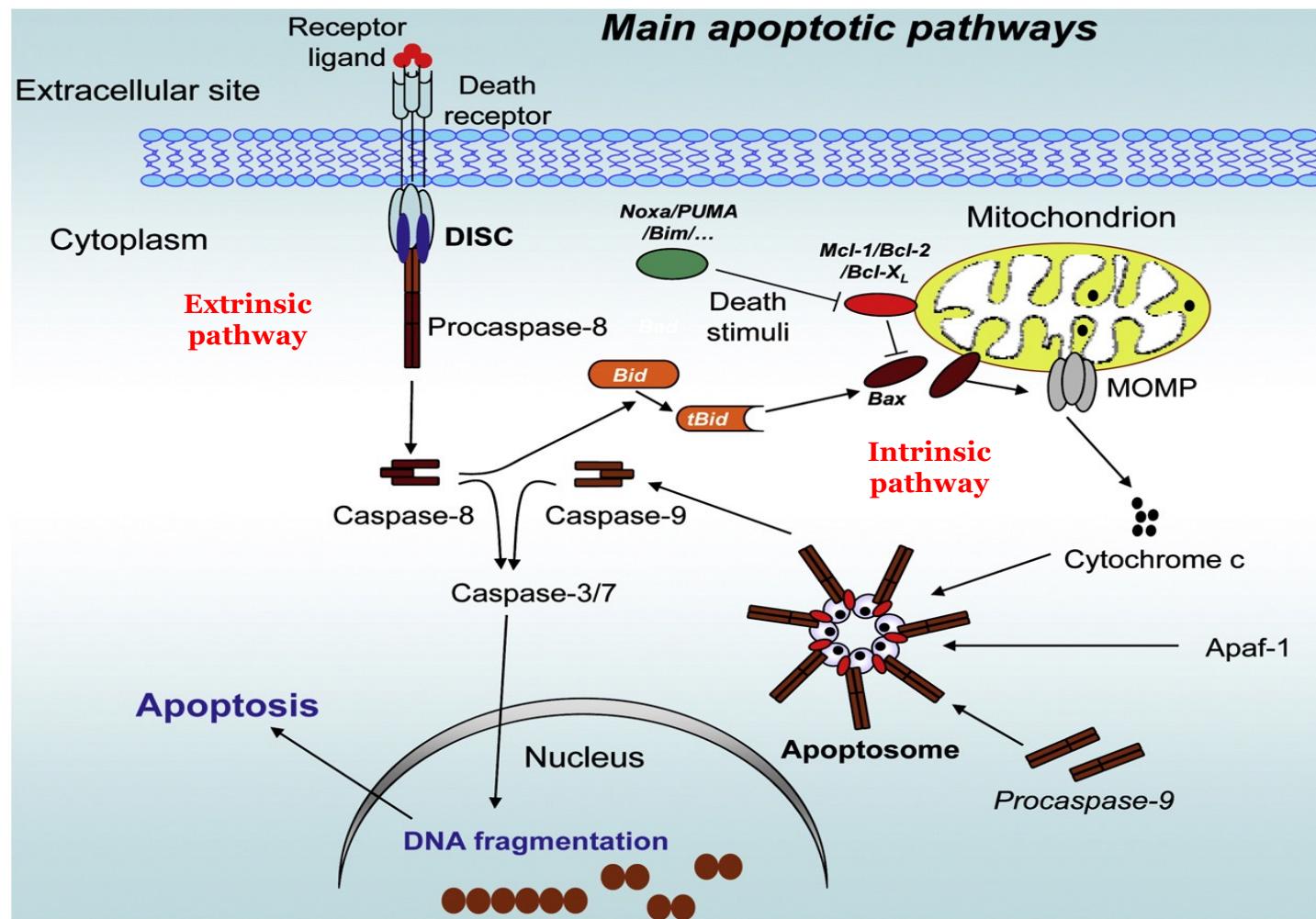


# **Production of Bcl-2 proteins involved in regulation of mitochondrial apoptosis**

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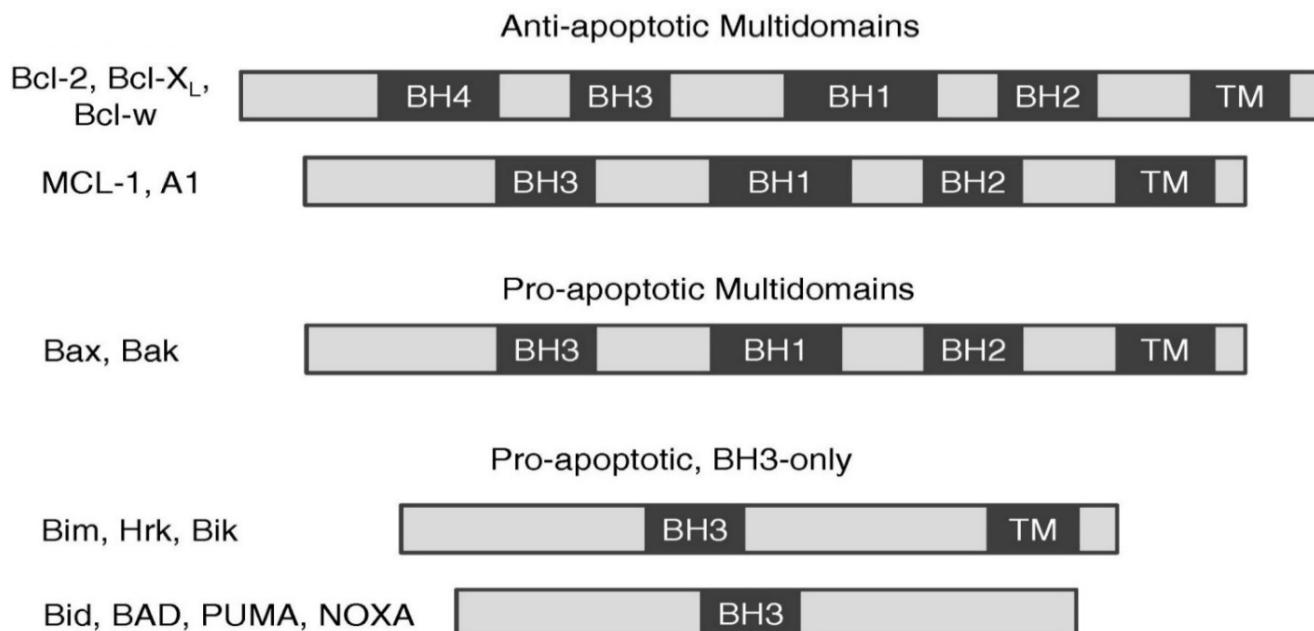
# Apoptosis is key for controlled removal of cells



\* Adapted from Nys K. et al., *Biochim Biophys Acta*, 2011

# The Bcl-2 protein family - regulators of the intrinsic pathway

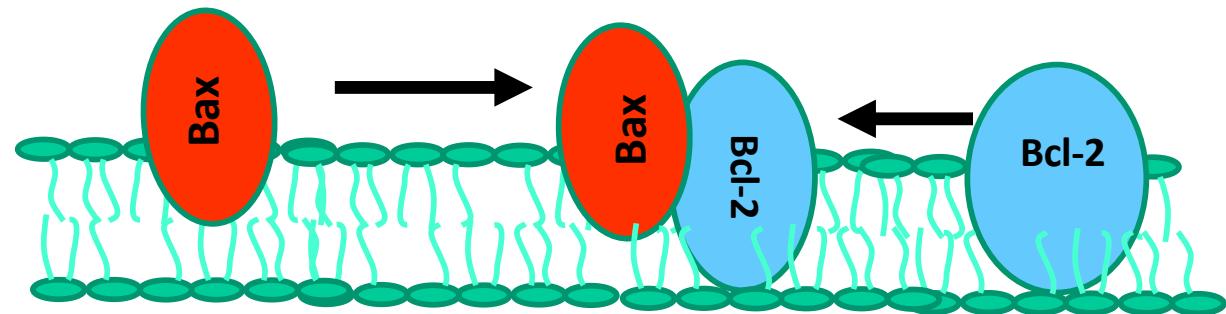
- Mainly  $\alpha$ -helical, with conserved BH-domains
- Anti- and pro-apoptotic classes, having opposing functions
- BH3-only proteins activate Bax and Bak



\* Adapted from Simpson C. D. et al., Cancer Lett, 2008

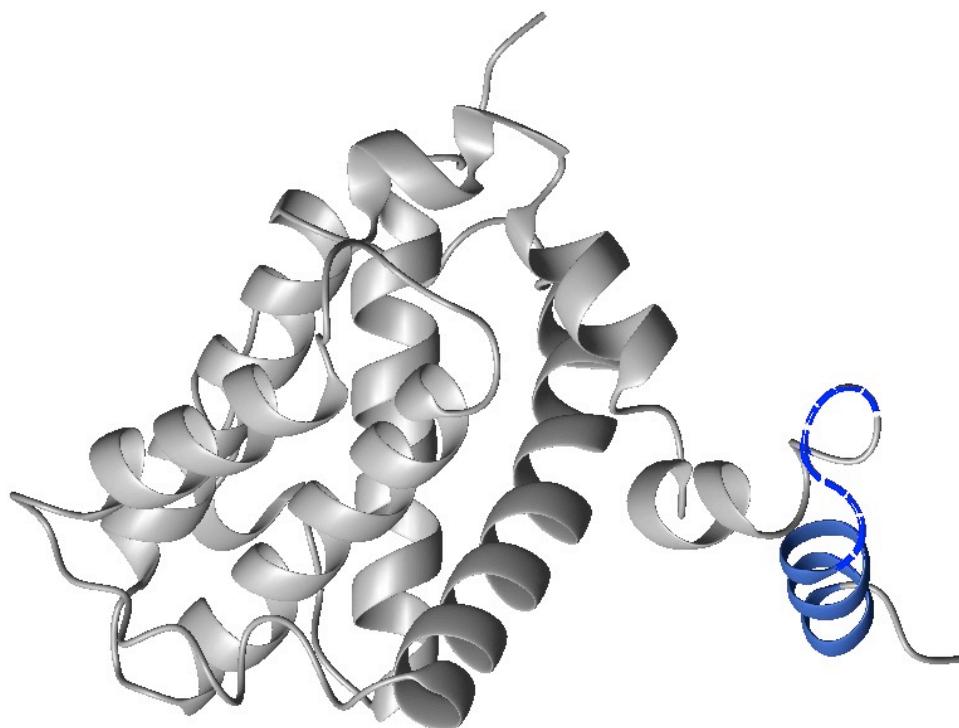
# How does the anti-apoptotic Bcl-2 protein function at the membrane level?

**Bcl-2 is a family of proteins that dictate the fate of the cell by a fine-tuned balance between anti- (Bcl-2) and pro-apoptotic (Bax) proteins that meet at the mitochondrial outer membrane.**



- Bcl-2 is an integral 26.5 kDa membrane protein
- Bcl-2 location not known
- Bcl-2 structure not known
- Bcl-2/Bax complex structure not known

# The anti-apoptotic membrane protein Bcl-2



**NMR structure of Bcl-xL. The C-terminal membrane anchor is shown in blue\***

\* A. Antignani, et al. J Mol Biol. 2015 Jul 3;427(13):2262-70, PDB ID: 1BXL

## Our aims

- **Solve the structure of the full-length anti-apoptotic membrane protein Bcl-2.**
- **Learn how the protein functions in the mitochondrial outer membrane.**
- **Study Bcl-2 together with Bax as it prevents it from forming membrane pores and trigger its cell-death inducing mechanism.**
- **Milligram amounts of protein is necessary when doing NMR and structural work.**

# Development of an expression method for Bcl-2 in *E. coli*

Over-expression in pET-15b at 37 C  
Grow BL21(DE3) Rosetta cells overnight in M9 media\*

Resuspend in Tris buffer at pH 8.0

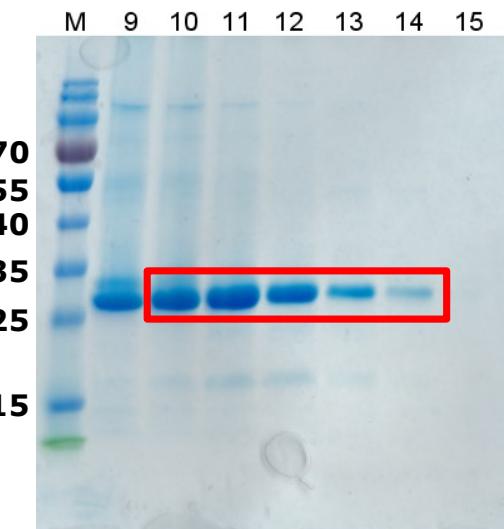
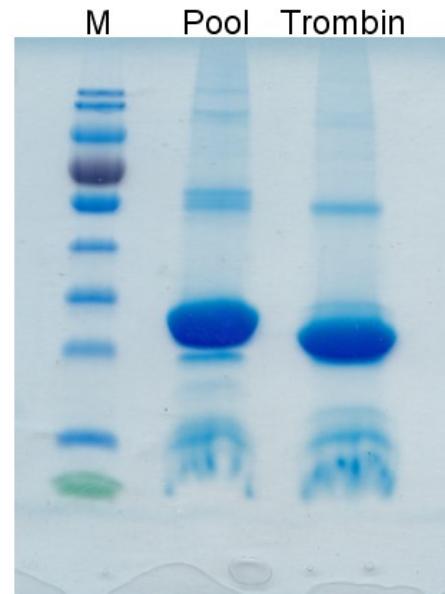
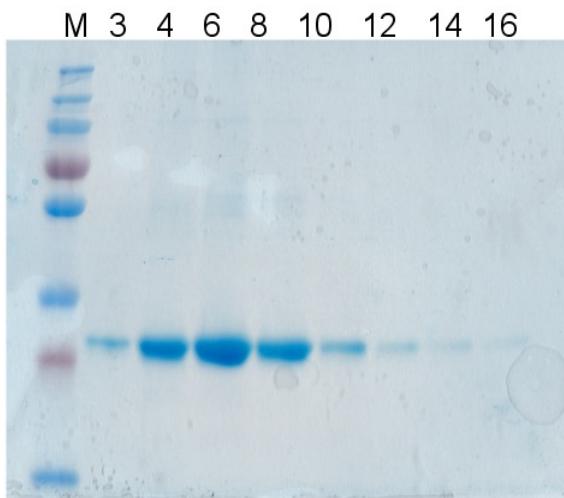
Solubilize pellet in 7 M urea, 0.6% sodium lauroyl sarcosinate,  
20 mM Tris, 200 mM NaCl, 10% (v/v) glycerol, pH 8.0

Dialysis against 20 mM Tris, 0.2 % Brij-35, 200 mM NaCl,  
10% glycerol, pH 7.8

IMAC purification with detergent exchange into DPC and  
removal of His-tag (thrombin)

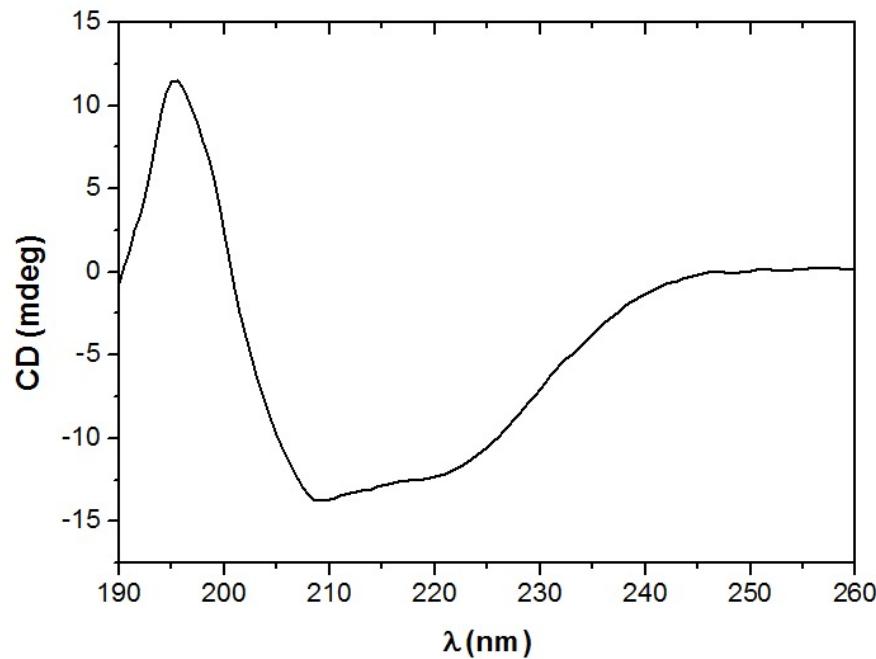
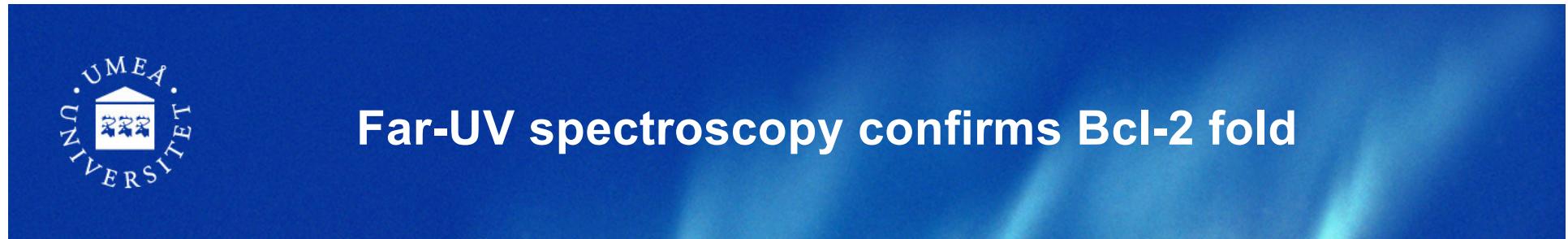
Gel filtration

# Expression and purification of Bcl-2 in *E. coli*.

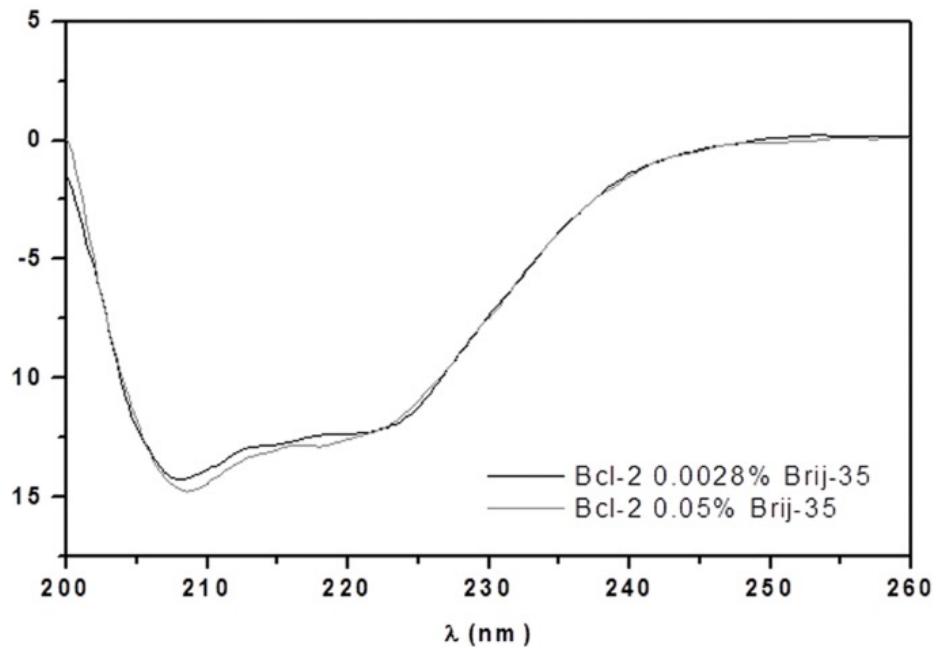


Bcl-2

Expression of Bcl-2 can now be achieved in LB or M9 media to yield  $^2\text{H}$ ,  $^{15}\text{N}$  or  $^{13}\text{C}/^{15}\text{N}$ -labeled protein



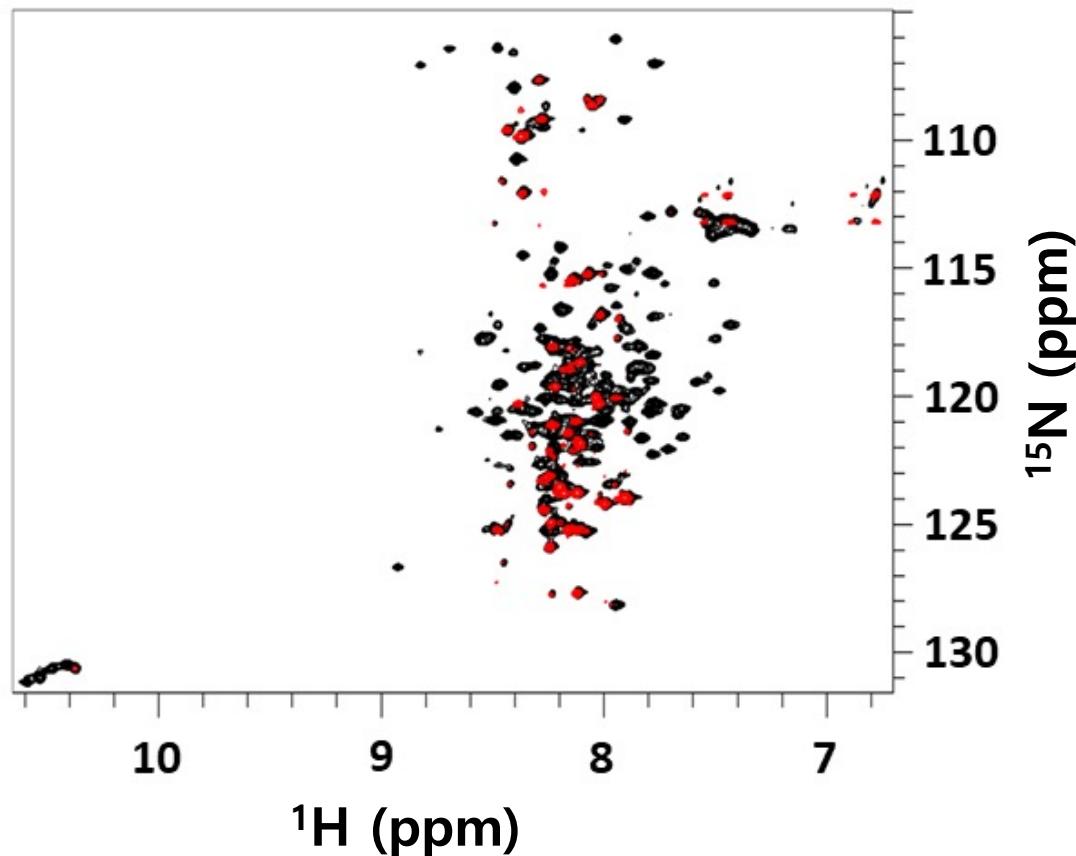
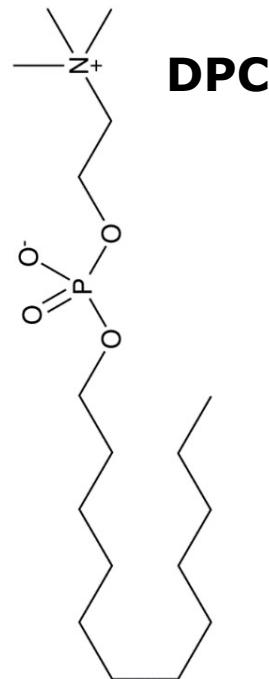
5  $\mu$ M His-tagged Bcl-2  
in 25 mM NaP, 20 mM KCl, 0.05 %  
Brij-35, 200  $\mu$ M DTT, pH 7.4, 25 C  
Expressed recombinantly in *E. coli*



5  $\mu$ M Bcl-2, in 25 mM NaP, pH 7.4,  
50 mM NaCl, 25 C  
Expressed previously using a cell-free  
expression system\*

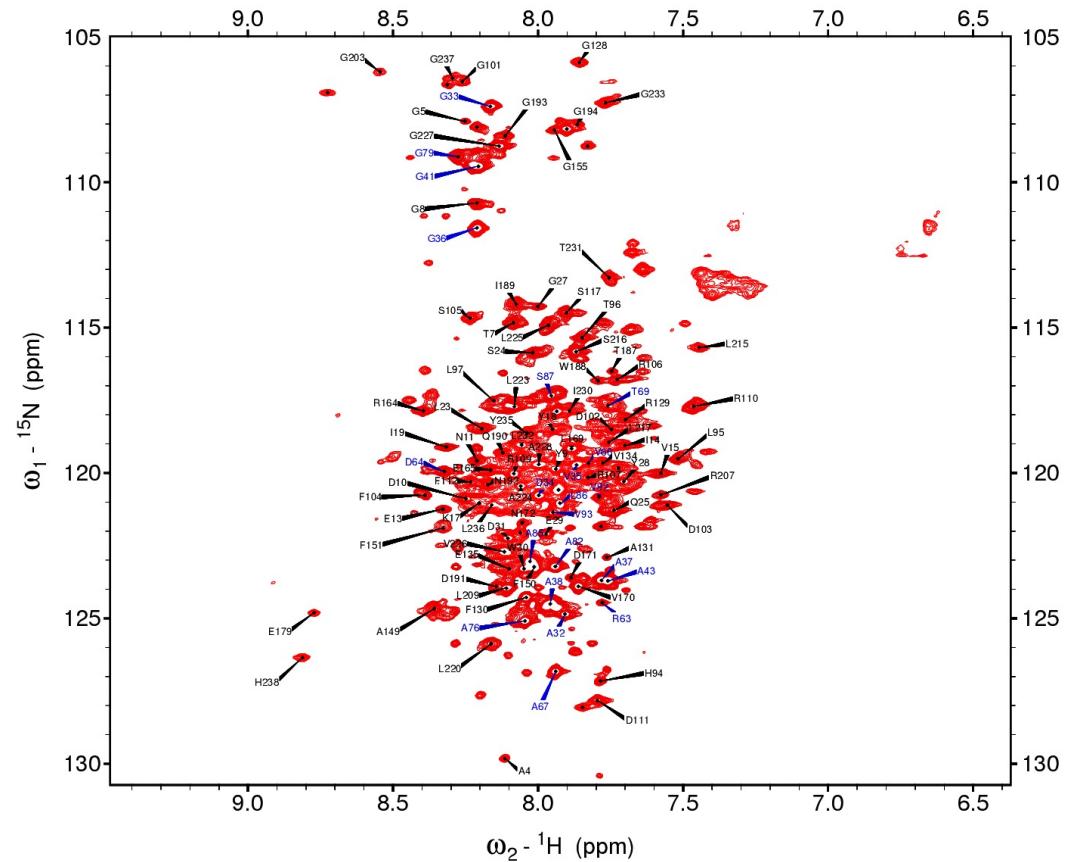
\* G. Gröbner et al. PLoS One. 2013; 8(4): e61452.

## Bcl-2 in DPC Micelles: Identification of flexible loop region residues



- We can identify the **free loop region residues** from  $T_2$  experiments with longer  $T_2$  delay  $\sim 410\text{ms}$
- Those residues must be outside the membrane and micellar core region
- Which residues are involved?

# Sequence-specific assignment of flexible loop region residues by NMR



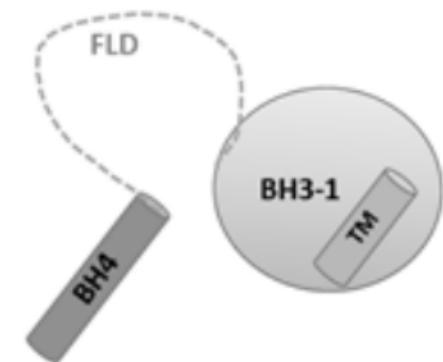
- All loop residues can be identified in the  $^1\text{H}$ - $^{15}\text{N}$  TROSY-HSQC spectrum.
- Residue-specific changes can be observed upon regulative activities such as monitoring of ligand/drug binding or interaction with other Bcl-2 proteins.
- Full-length spectrum of Bcl-2 is however crowded with several overlaps.

# “Divide-and-conquer” strategy to improve the spectral resolution of Bcl-2

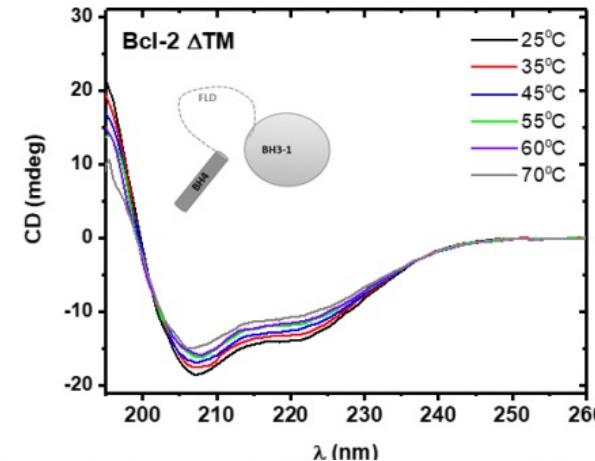
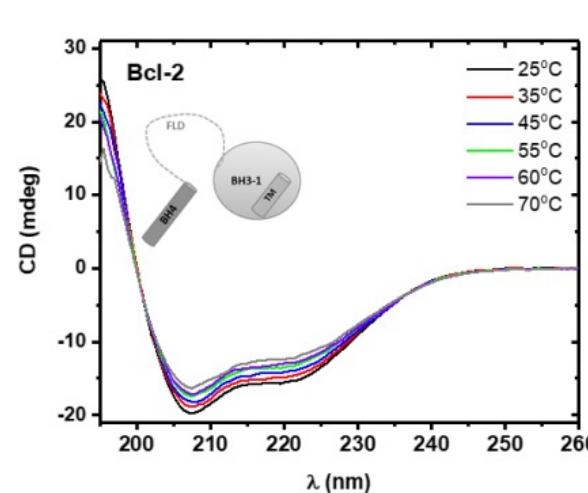
- Truncate the protein at various positions to overcome problems with overlap.
- Use several spectra together to help with residue assignments and/or to define functional regions within the protein.
- Use this strategy to aid structural determination.

## Strategy

- Make shorter Bcl-2 constructs by adding stop codons, terminating the sequence before its trans-membrane helix.
- Other constructs made to omit the BH4 domain and the FLD.

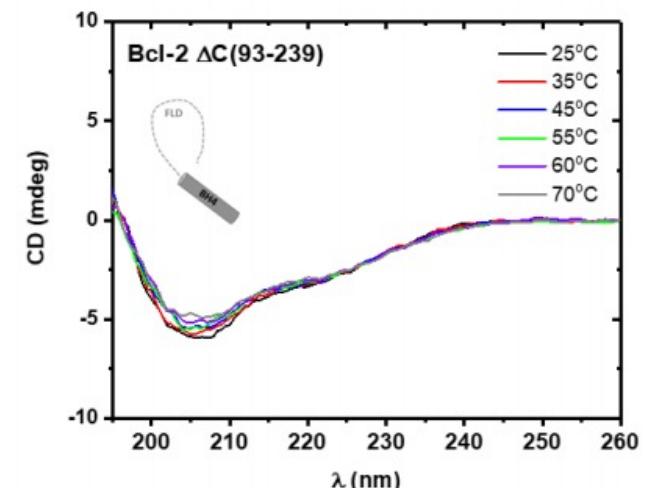
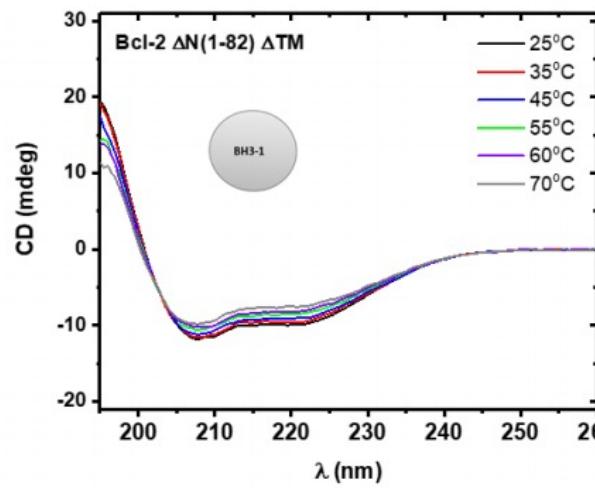
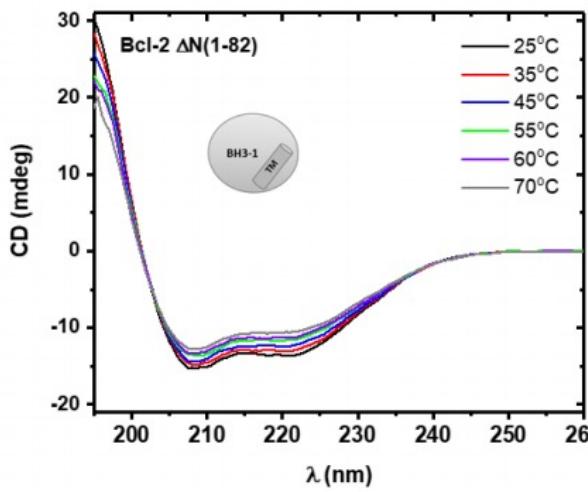


# Far-UV CD of Bcl-2 and truncated variants in DPC micelles



Partially soluble

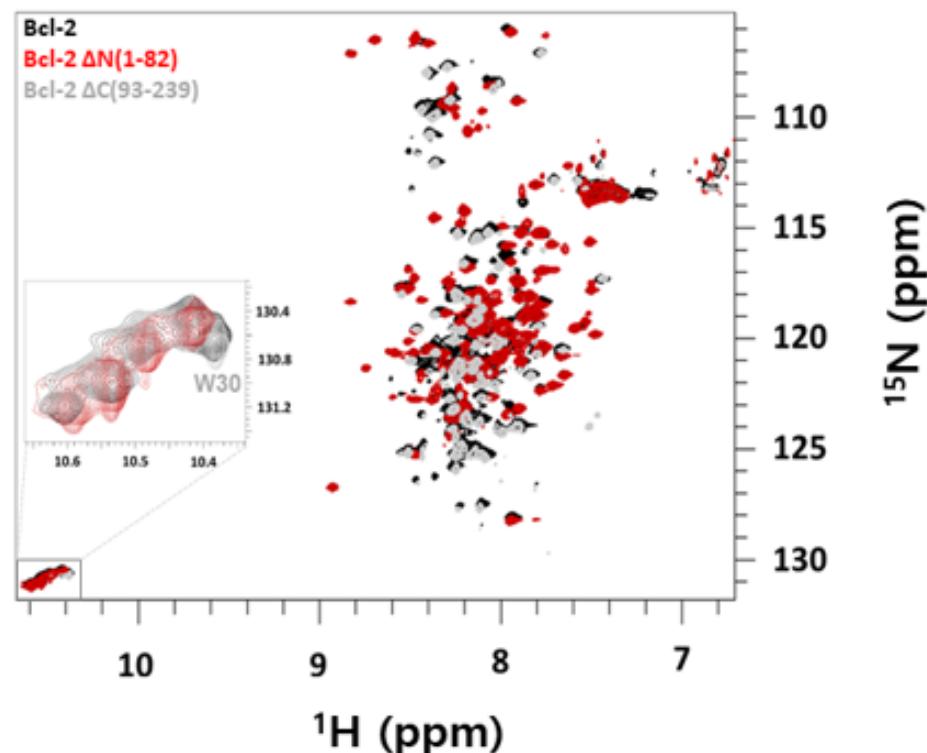
Recorded at pH 6.0 in DPC



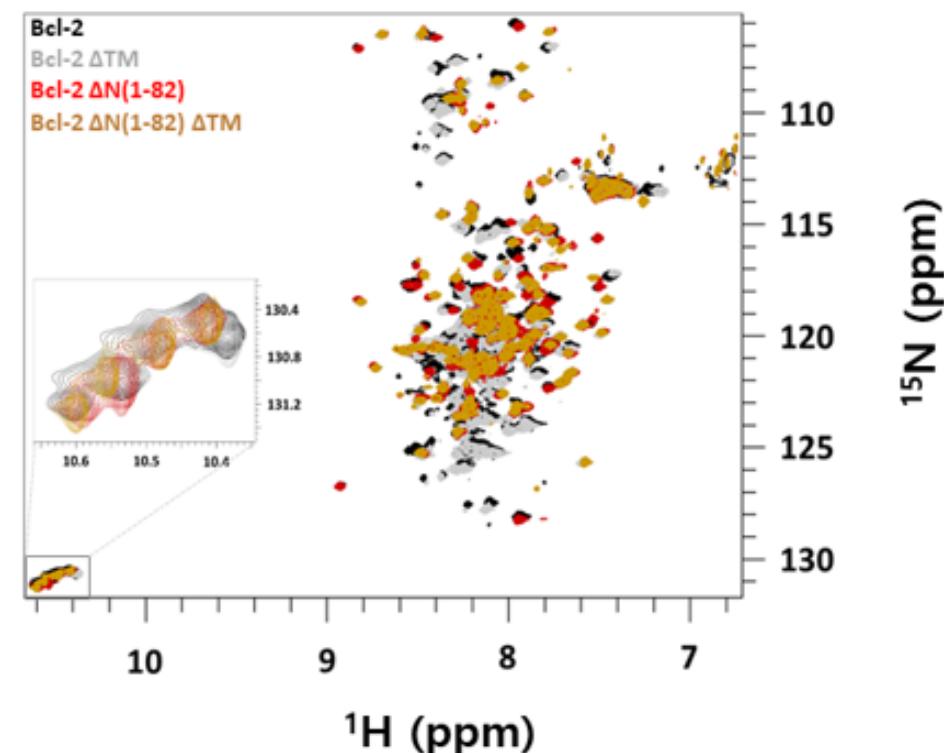


# Characterization of truncated Bcl-2 constructs

**A**



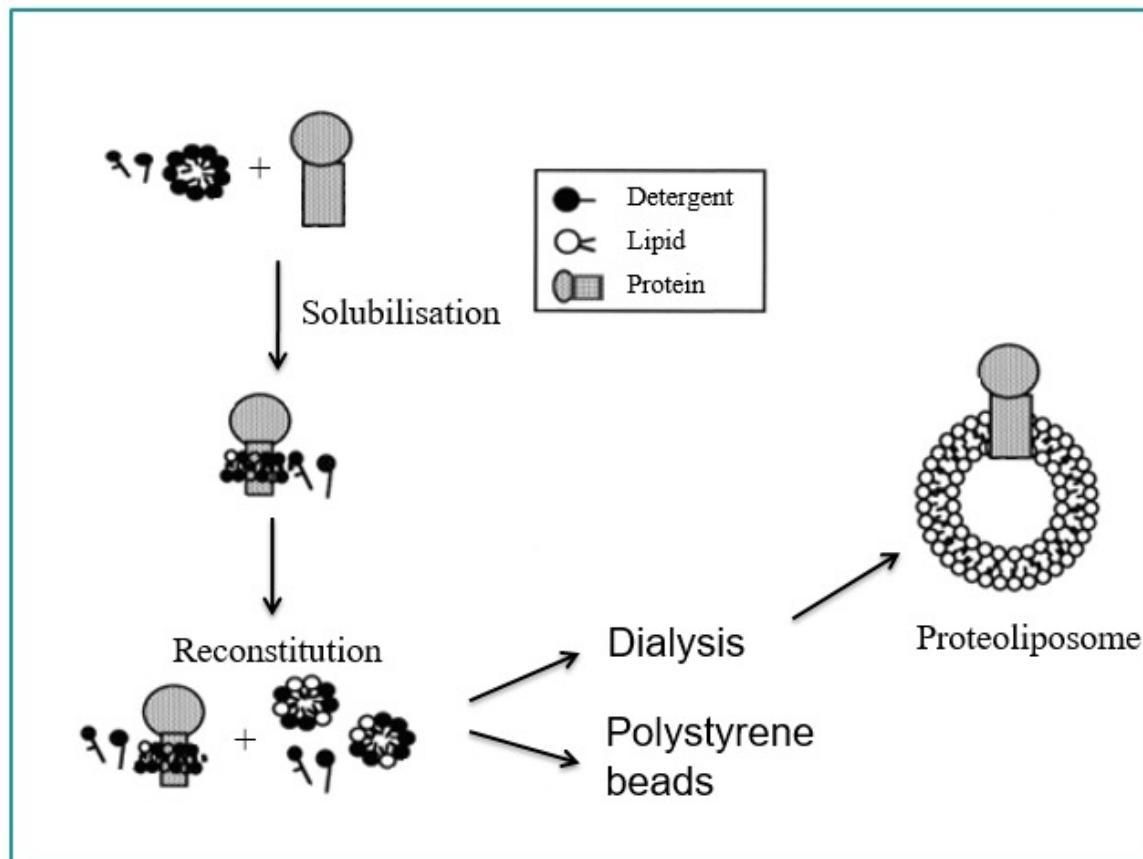
**B**



- Improved spectral resolution
- Identification of some residues by simple comparison

# Reconstitution of Bcl-2 into DMPC vesicles

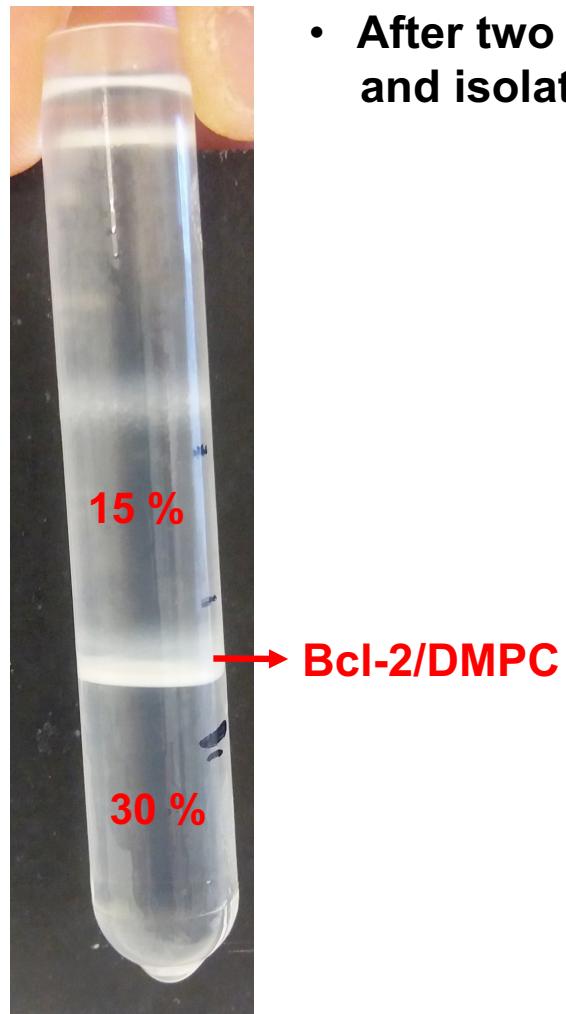
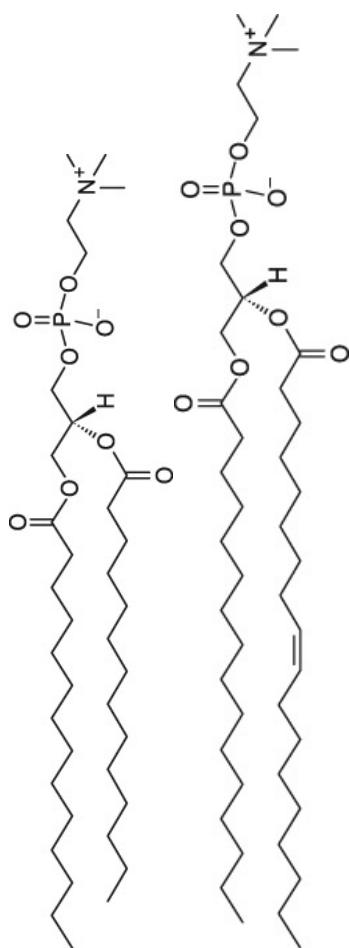
**Incorporation of Bcl-2 into vesicles, for studies under native-like conditions**



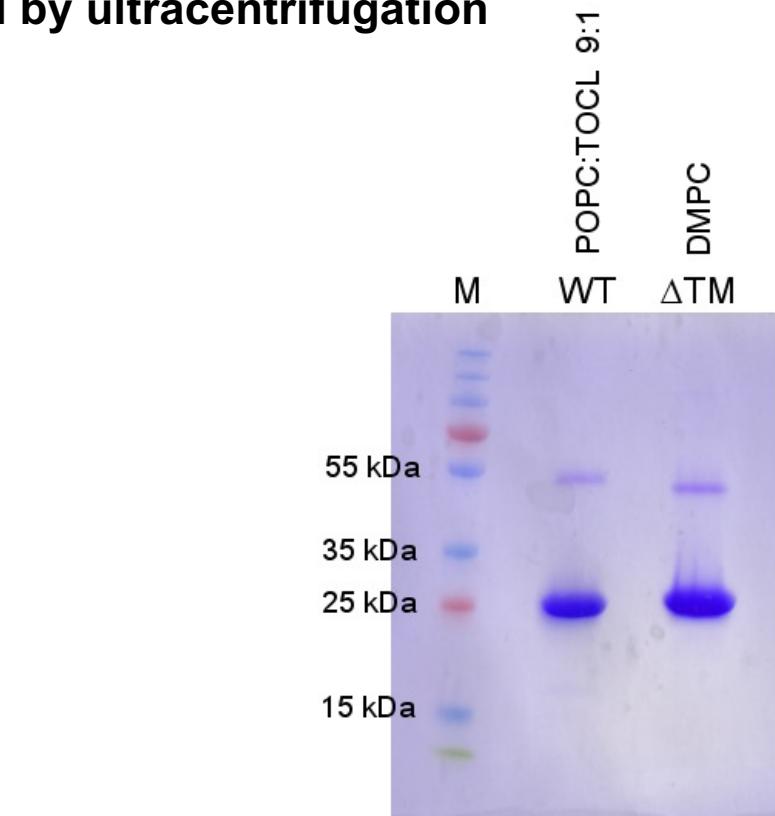
Adapted from Rigaud J.-L., Braz J Med Res, 2002

# Reconstitution of Bcl-2 into proteoliposomes

DMPC POPC



- Removal of detergent with Bio-Beads SM-2 adsorbents at 4 C.
- After two days, proteoliposomes were obtained and isolated by ultracentrifugation



SDS-PAGE of reconstituted Bcl-2 in DMPC and POPC/TOCL liposomes



## Conclusions and future perspectives

- ✓ An alternative expression method for the human Bcl-2 membrane protein has been established, yielding milligram amounts of protein.
- ✓ Bcl-2 has been successfully incorporated into DMPC and POPC/CL vesicles.
- ✓ DPC is a robust detergent for structural and functional studies.
- ✓ Different shorter Bcl-2 constructs can be used together to form a full picture of the intact Bcl-2 membrane protein.



# Acknowledgements

## Group/NMR Facility

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- **Magnus Wolf-Watz**
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