

NMR study of viral IDPs

Tomáš Hošek,
CERM, Italy



Presentation overview

- IDPs & viruses
- Studied viral oncoproteins:
 - E7 from Human Papilloma virus 16
 - E1A from Adenovirus 2/5

Intrinsically Disordered Proteins

- function coded by short AA sequence (Short Linear Motifs)
- highly prevalent in eukaryotes
- also vastly exploited by viruses
- viral IDPs involved in cancer development

Dunker, A. K. (2001). Intrinsically disordered proteins. *Journal of Molecular Graphics and Modelling*, 19(1), 26–59.

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Davey, N. E., Travé, G., & Gibson, T. J. (2011). How viruses hijack cell regulation. *Trends in Biochemical Sciences*, 36(3), 159–169.

Uversky, V. N., & Dunker, A. K. (2008). Controlled chaos. *Science (New York, N.Y.)*, 322(5906), 1340–1341.

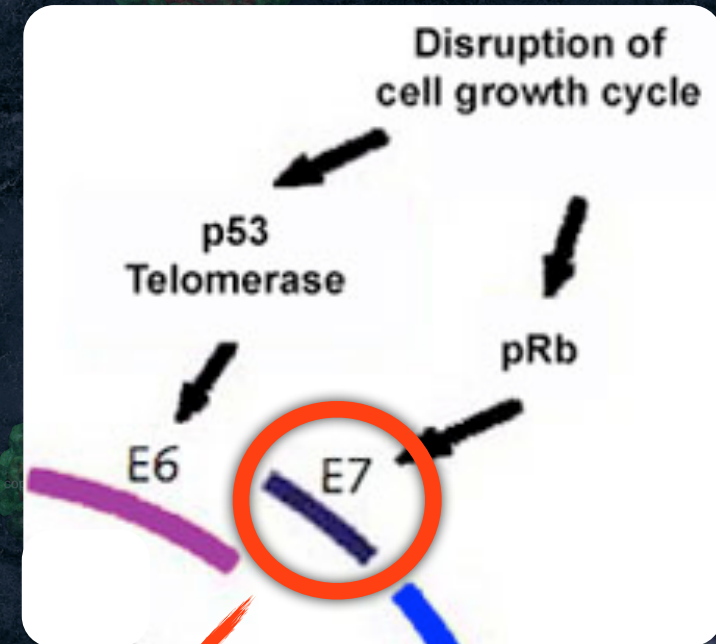
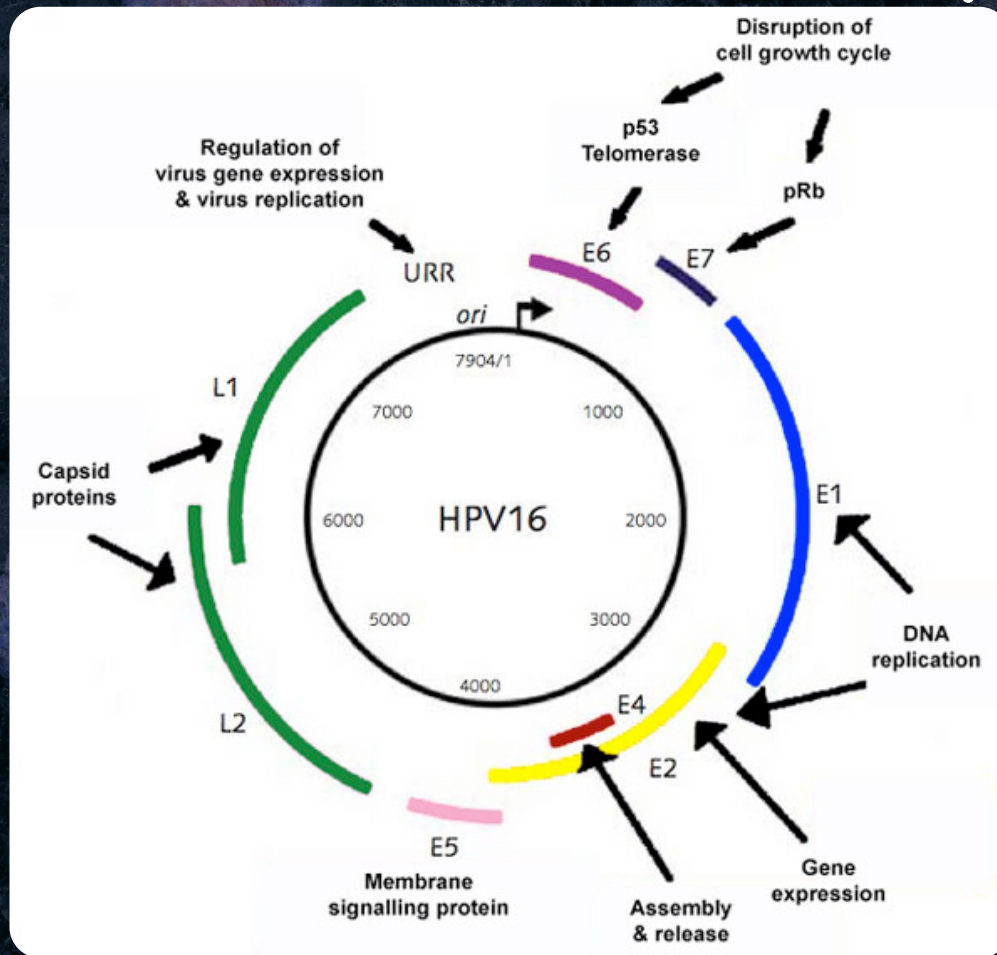


Human Papilloma viruses

- non-enveloped dsDNA viruses
- infect keratinocytes of skin or mucous membrane
- can cause cervix and throat cancer
- low-risk (HPV-6, HPV-11)
- moderate-risk (HPV-26, HPV-53)
- high-risk (HPV-16, HPV-18) = oncogenic

Uversky et al. (2006). Protein Intrinsic Disorder and Human Papillomaviruses: Increased Amount of Disorder in E6 and E7 Oncoproteins from High Risk HPVs. Journal of Proteome Research, 5(8), 1829-1842.

Human Papilloma virus

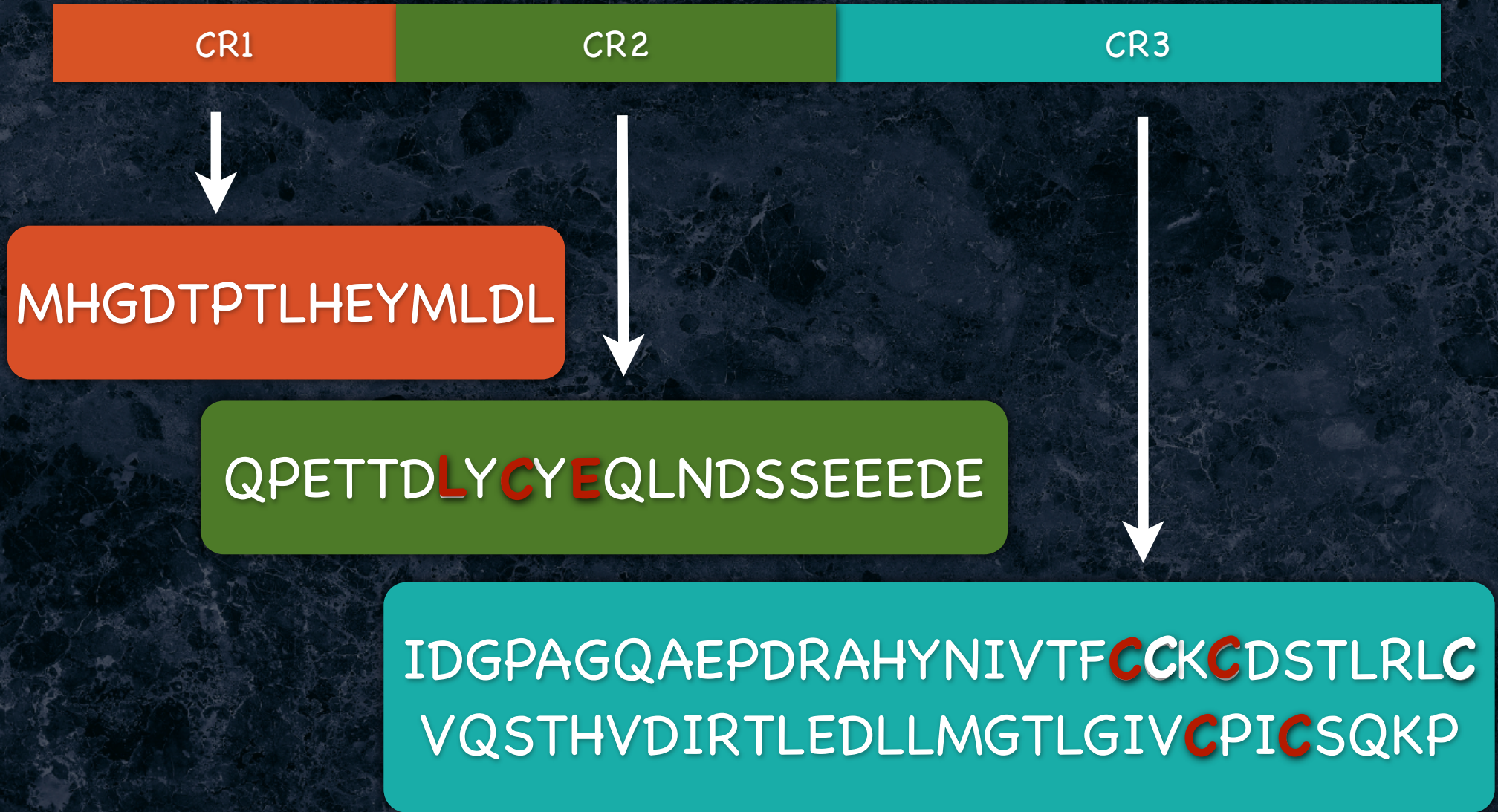


CR1

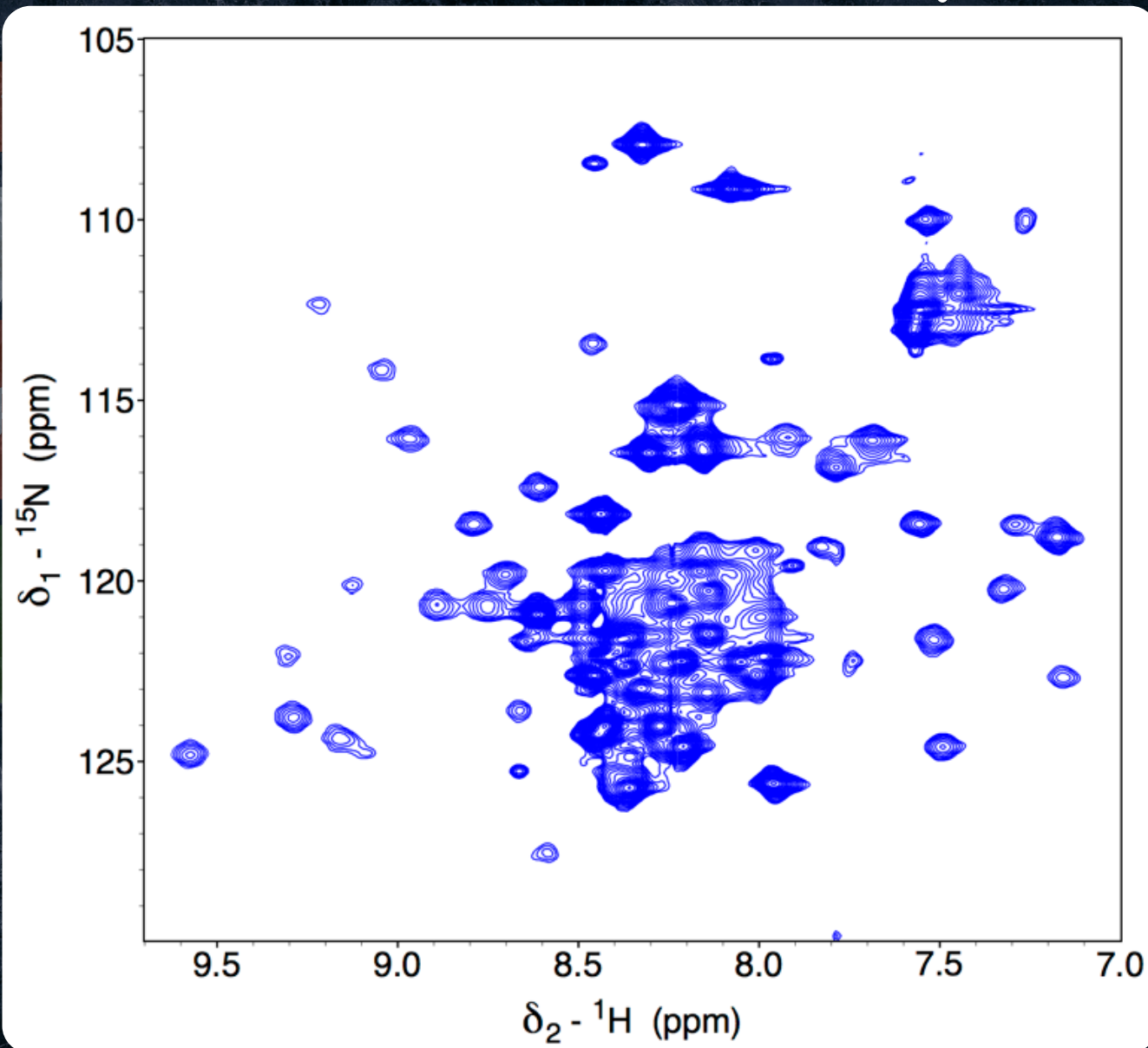
CR2

CR3

HPV 16 E7 oncoprotein



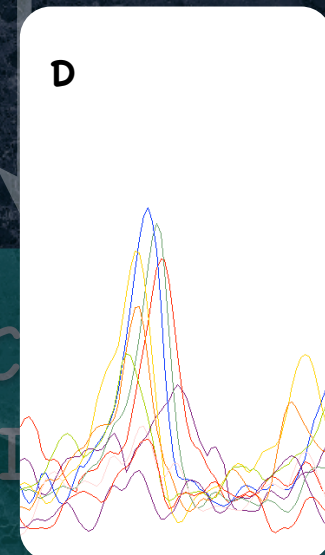
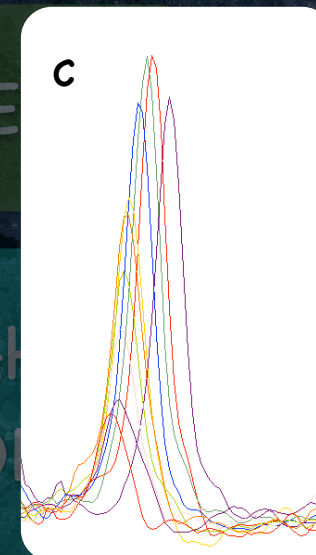
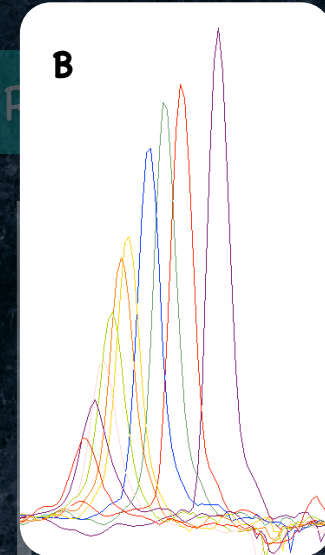
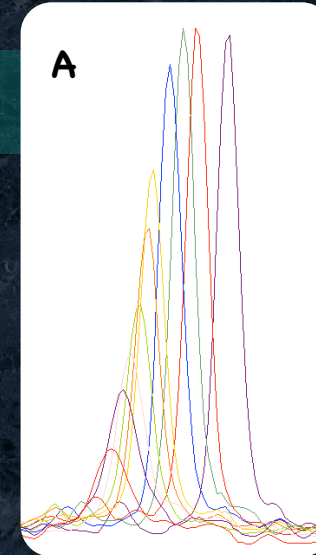
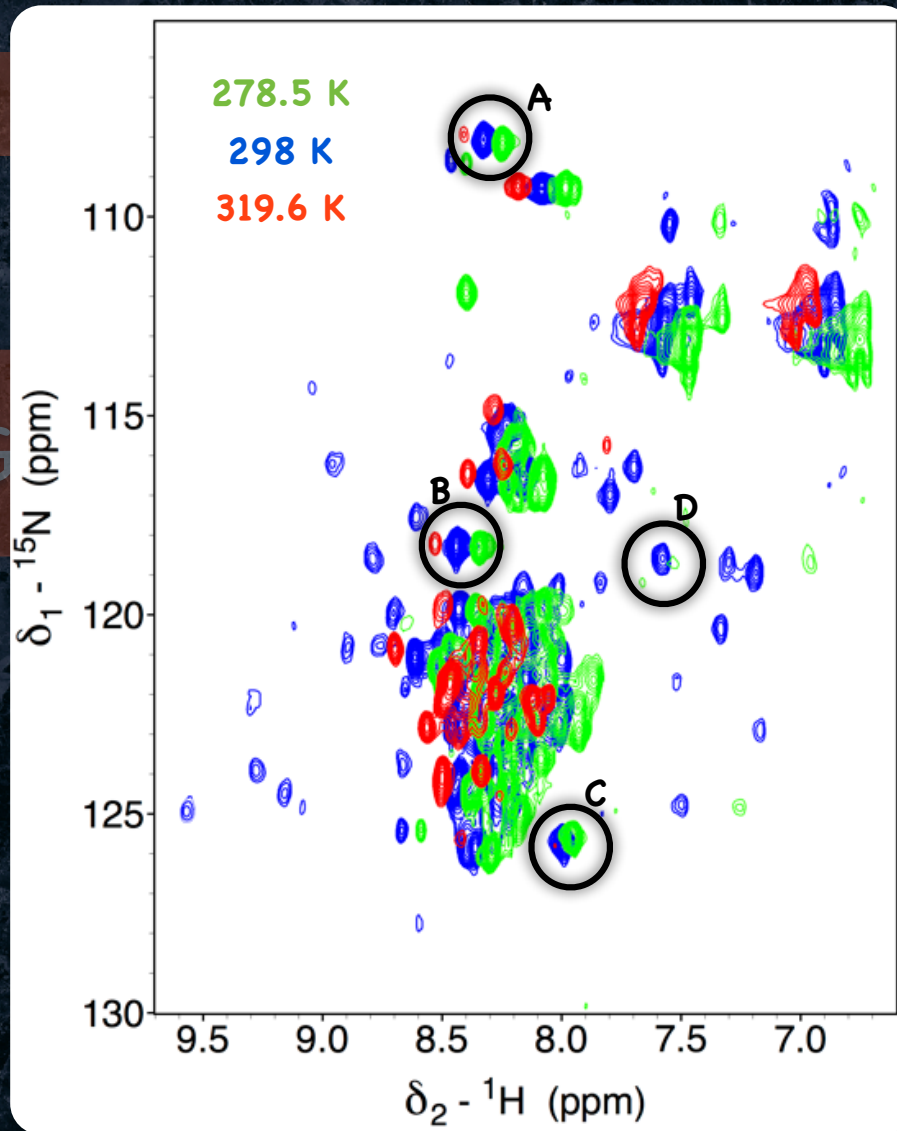
HPV 16 E7 oncoprotein



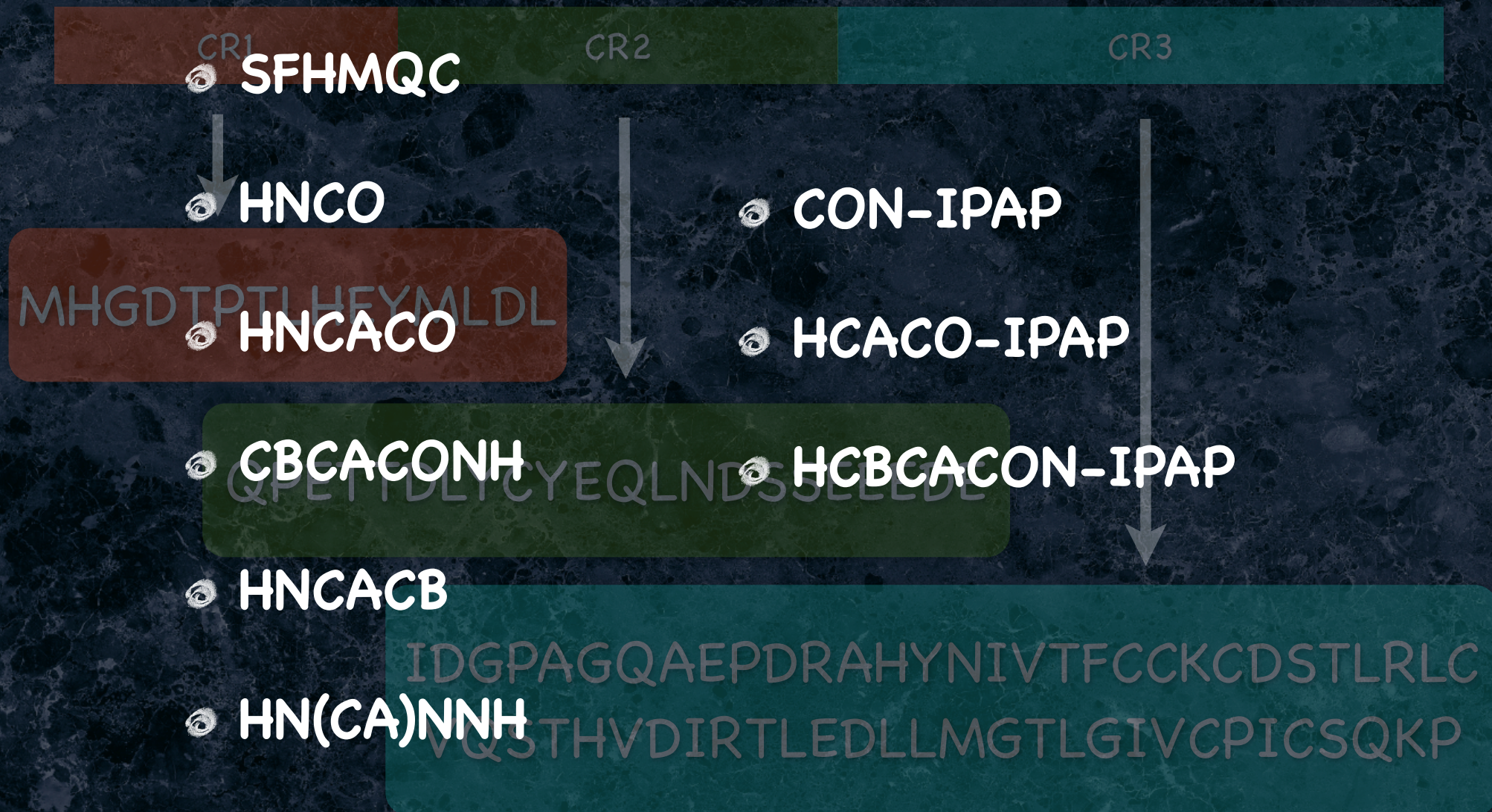
MHGDTR

CDSTLRRC
PICSQKP

Temperature dependence



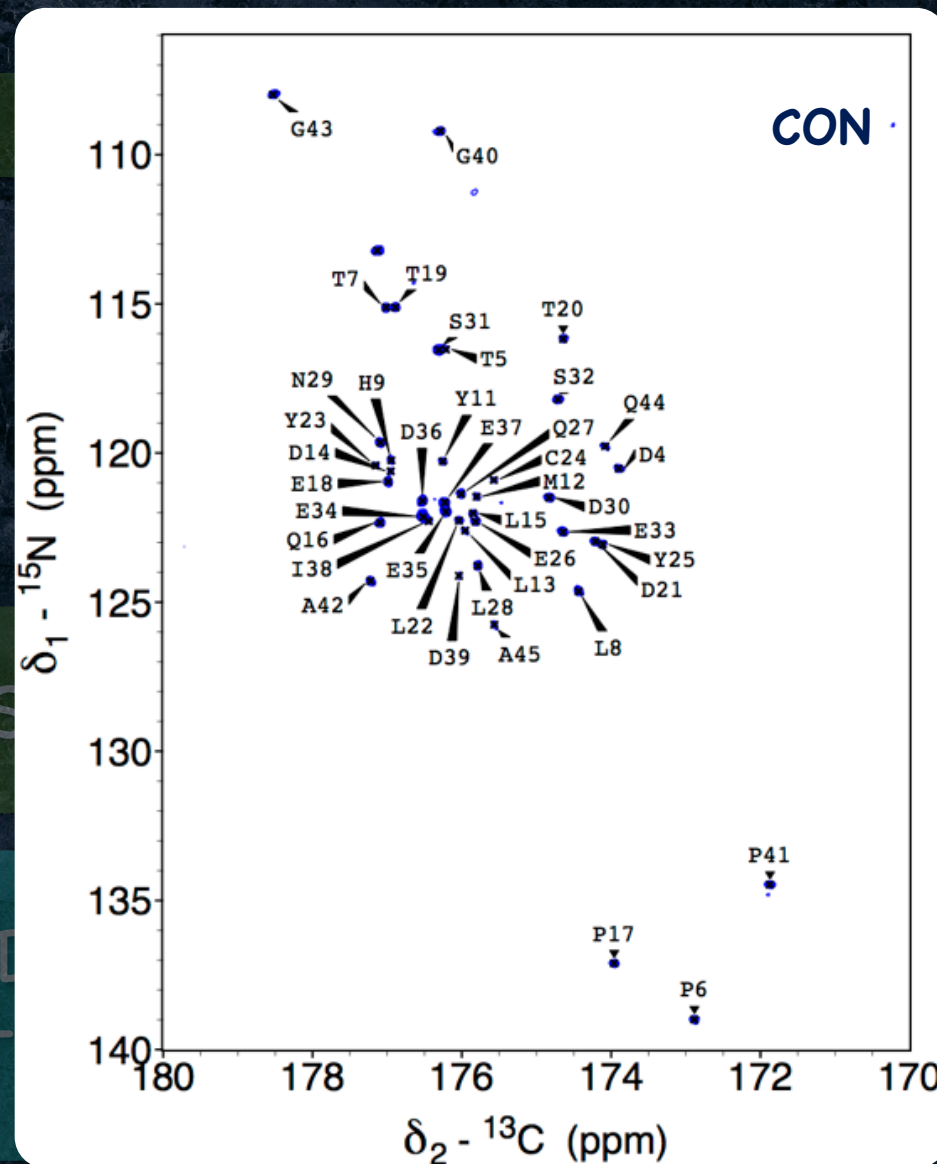
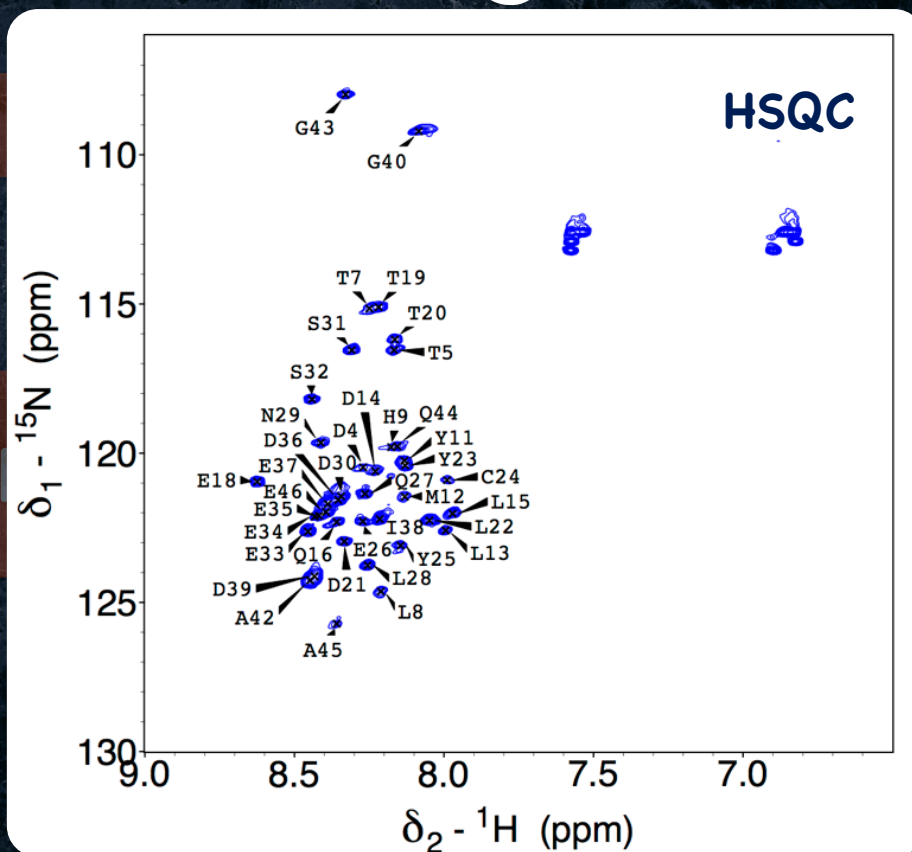
Assignment of HPV 16 E7



Assignment of HPV 16 E7

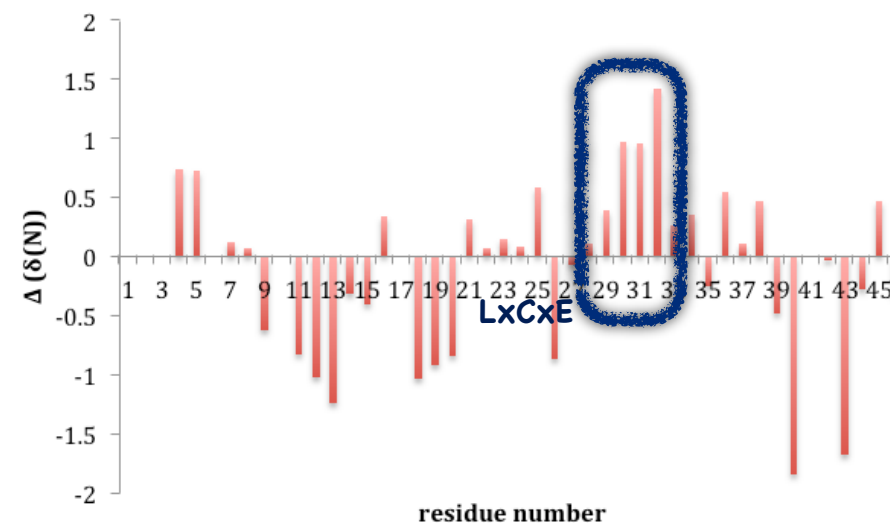
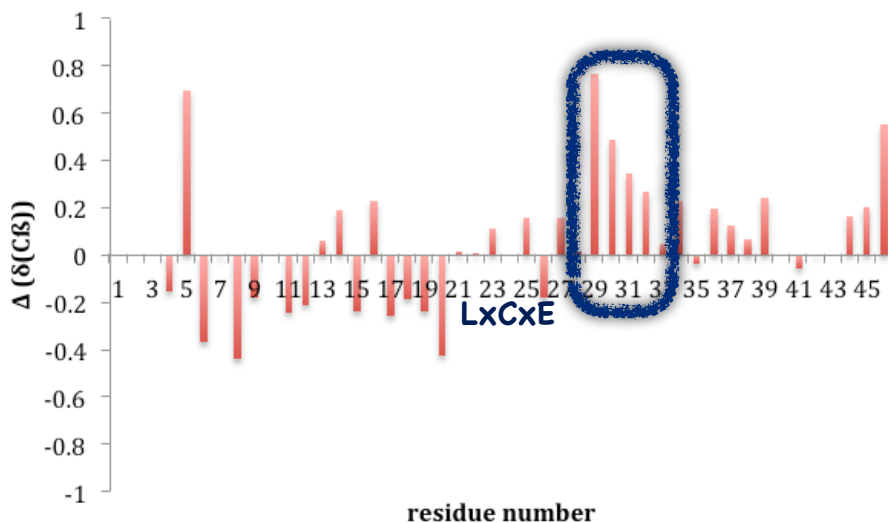
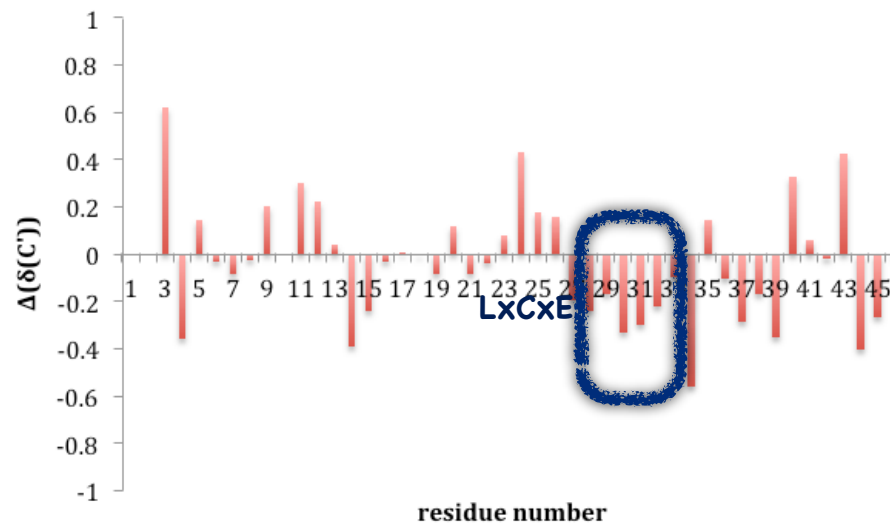
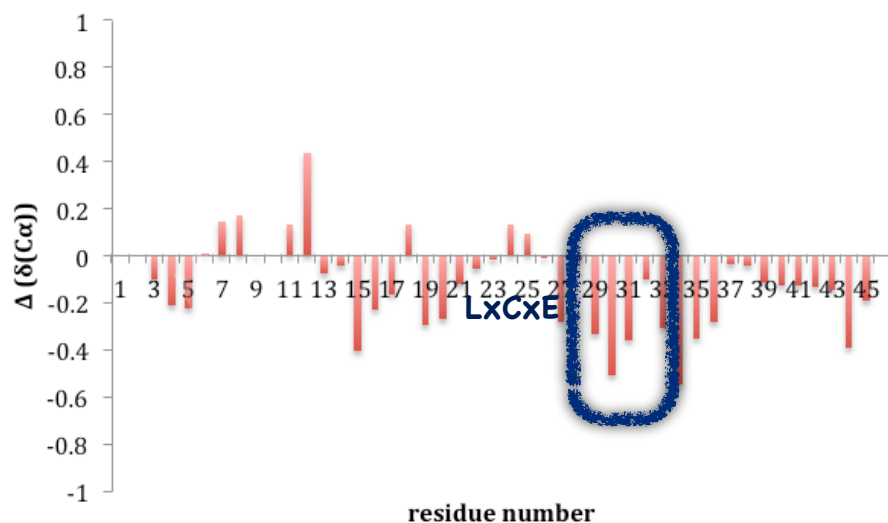


Assignment of HPV 16 E7



IDGPAGQAEPT
VQSTHVDIRT

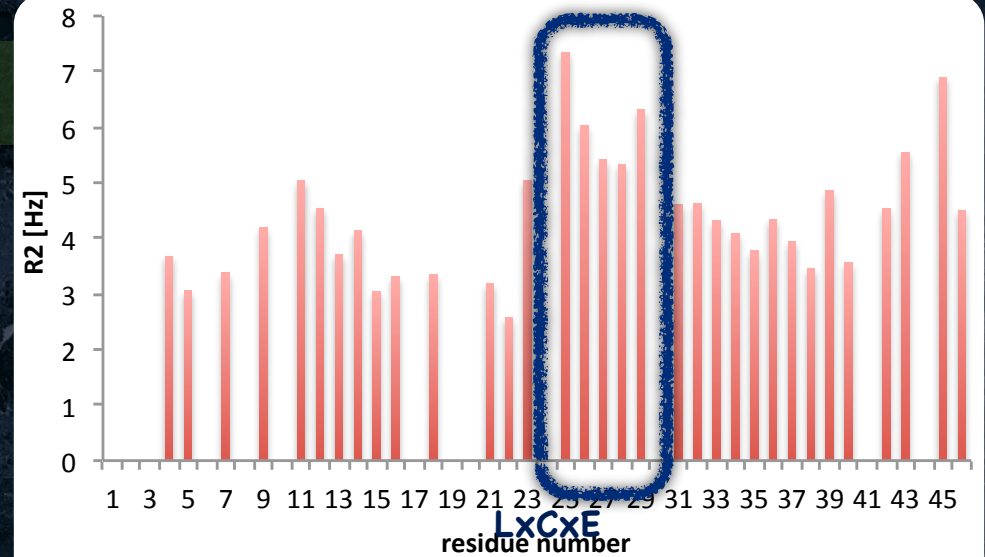
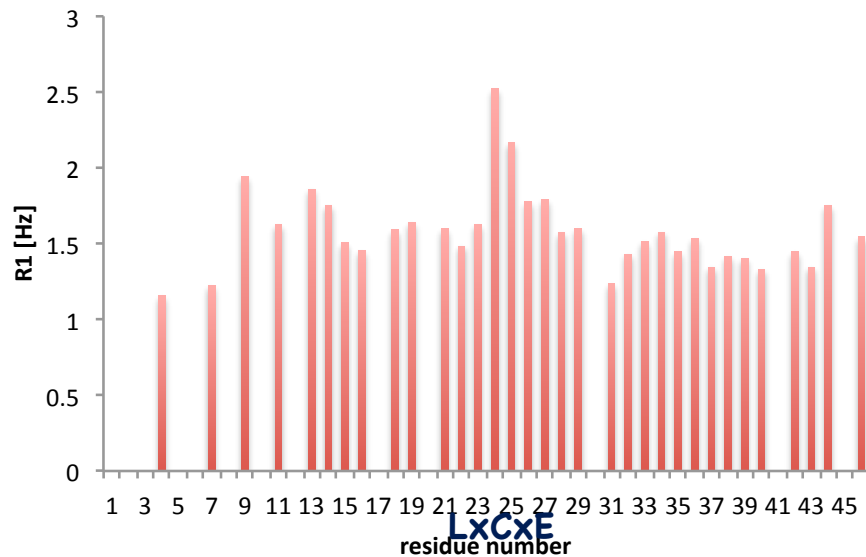
Chemical Shift Index



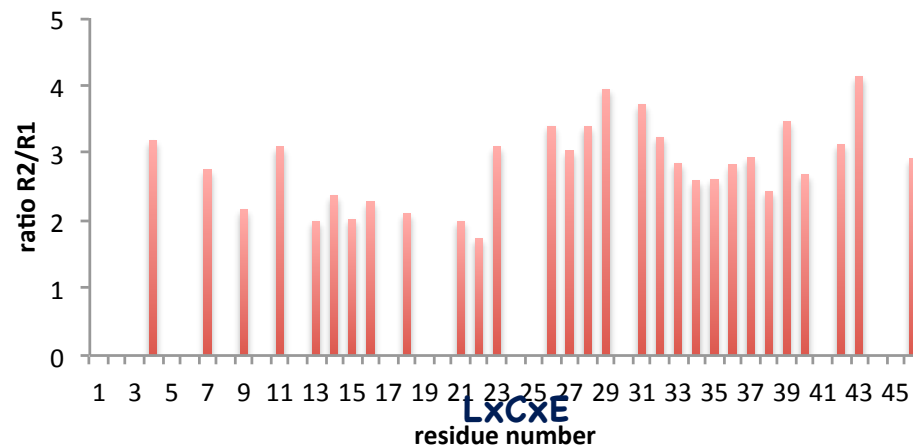
Lee, J., Russo, A. A., Pavletich (1998). Structure of Retinoblastoma tumor-suppressor pocket domain bound to a peptide from HPV E7. *Nature*, 391(6670), 859–865.

IDPbyNMR - High resolution tools to understand the functional role of protein intrinsic disorder - Project n. 264257

^{15}N relaxation experiments

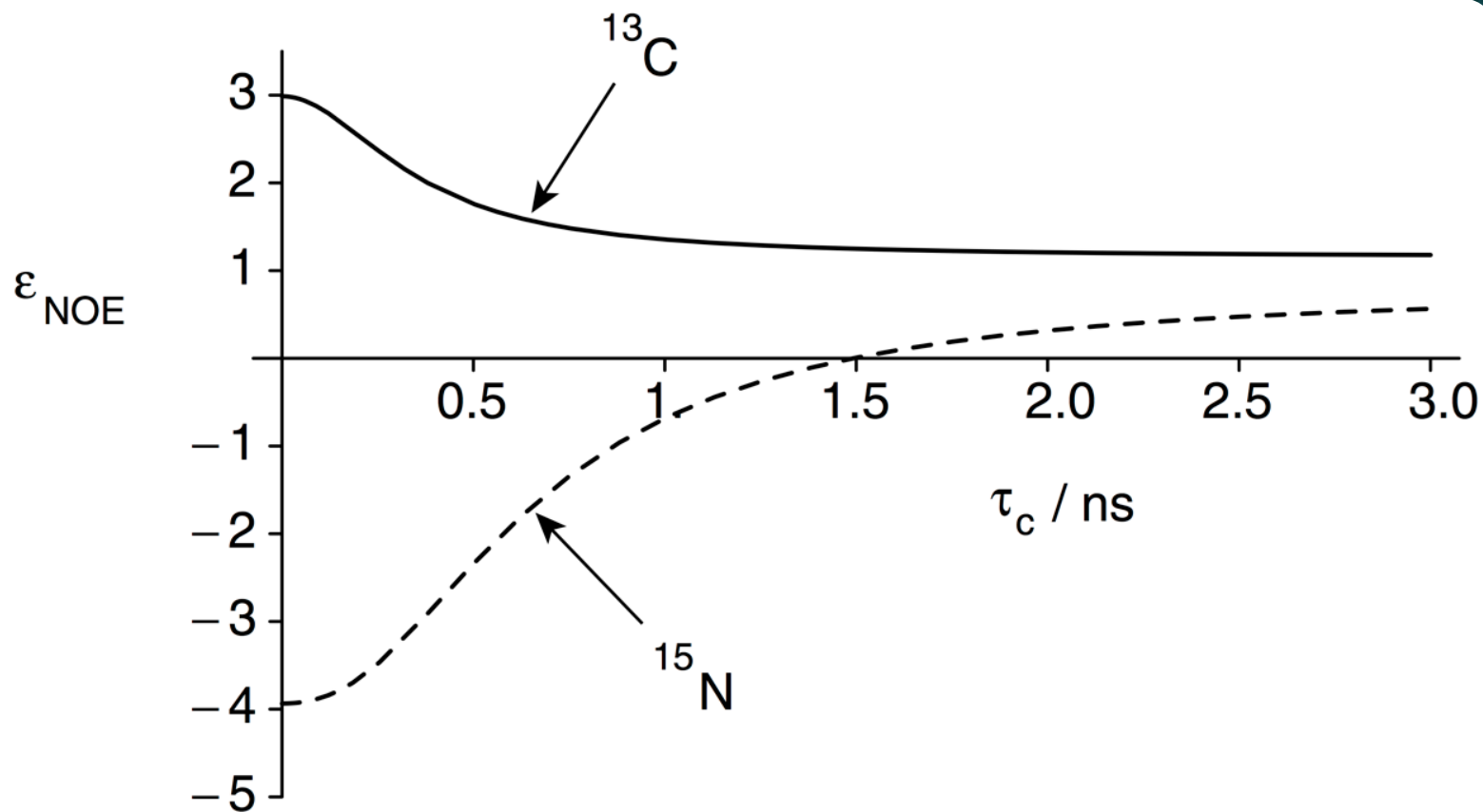


Correlation time



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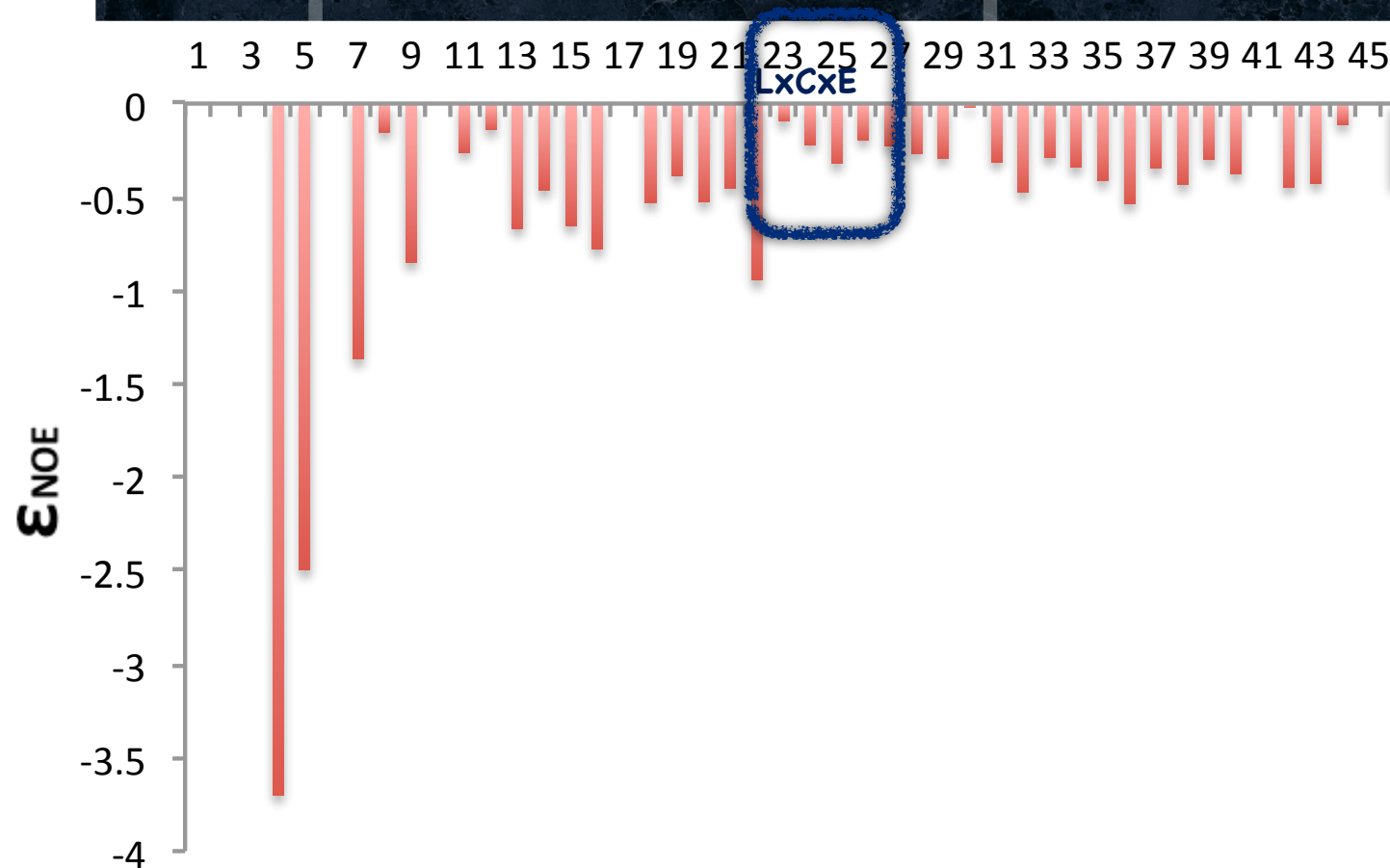
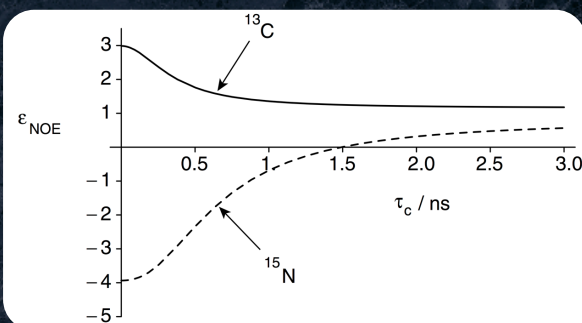
heteronuclear NOE



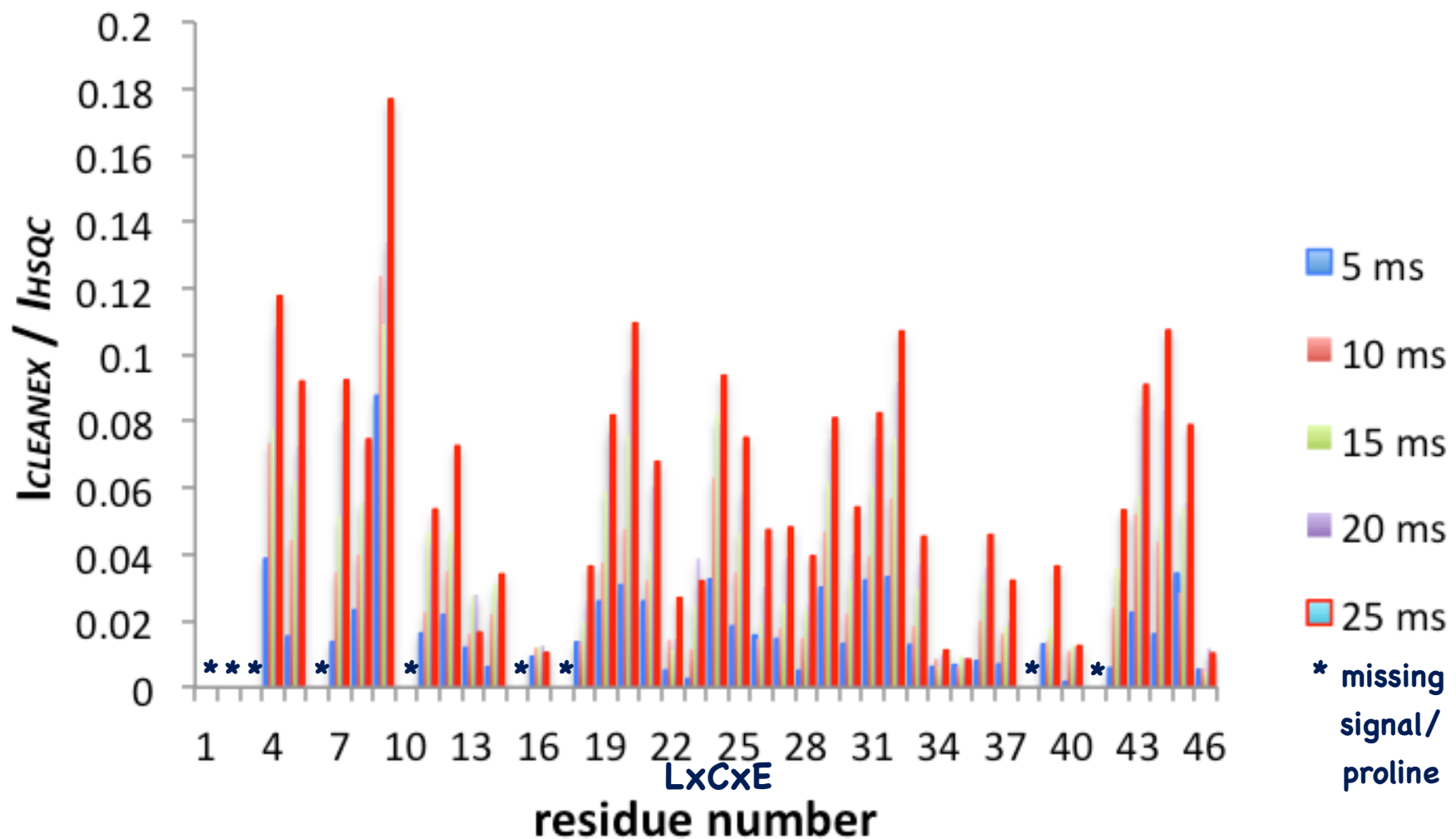
Malcolm H. Levitt, Spin Dynamics: Basics of Nuclear Magnetic Resonance - 2nd ed., Chichester: John Wiley & Sons Ltd., 2009

IDPbyNMR - High resolution tools to understand the functional role of protein intrinsic disorder - Project n. 264257

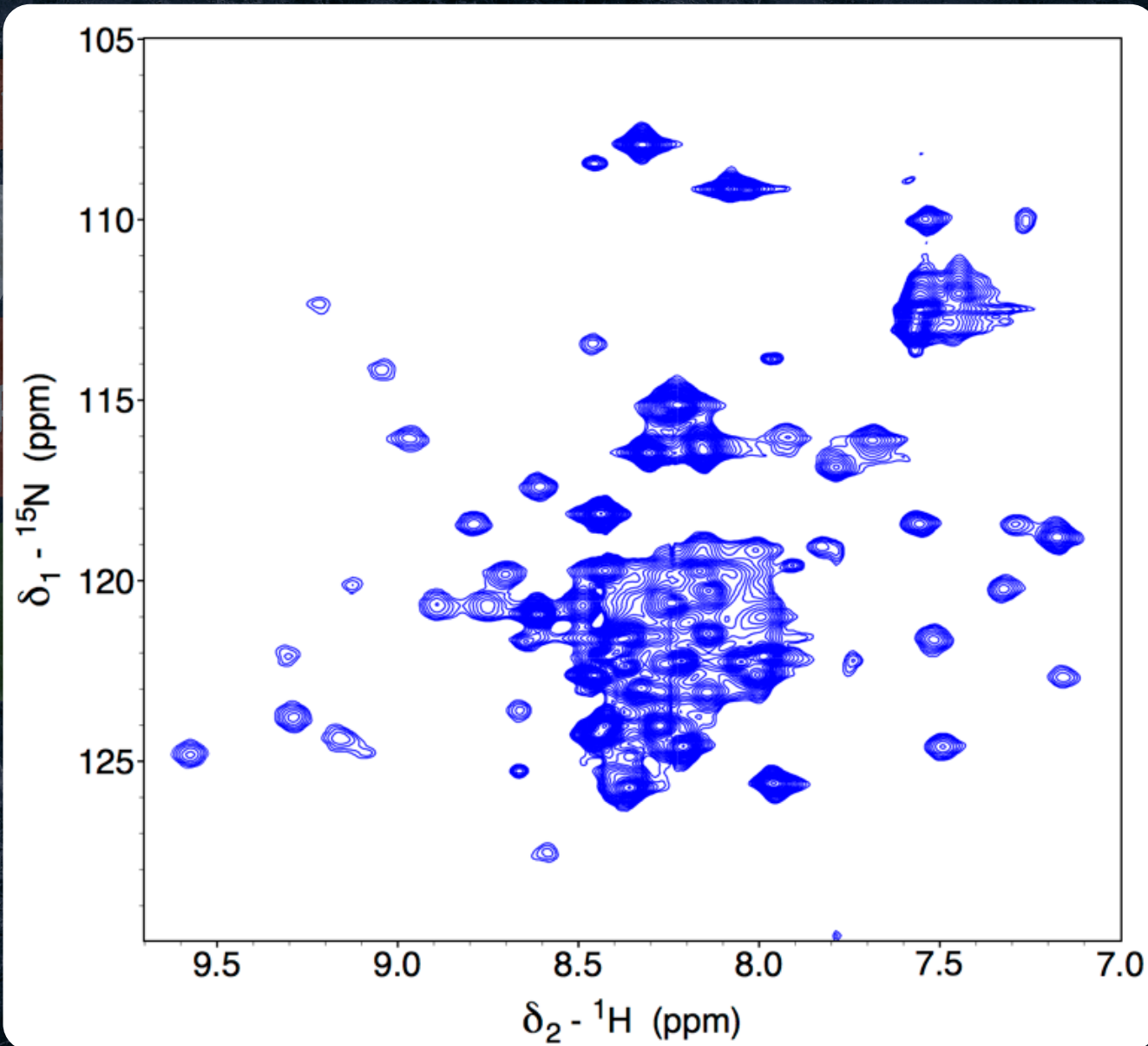
^1H - ^{15}N heteronuclear NOE



HN-H₂O exchange rates



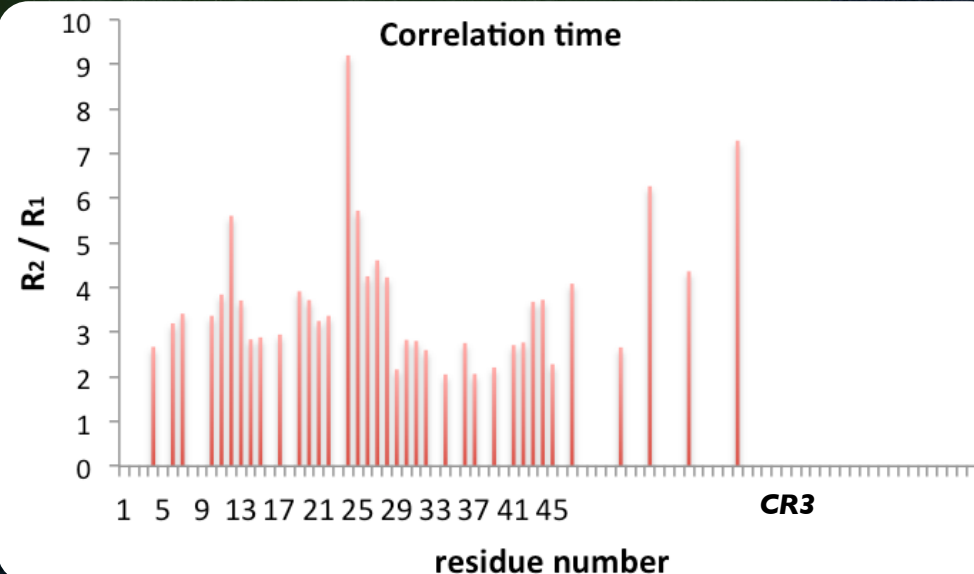
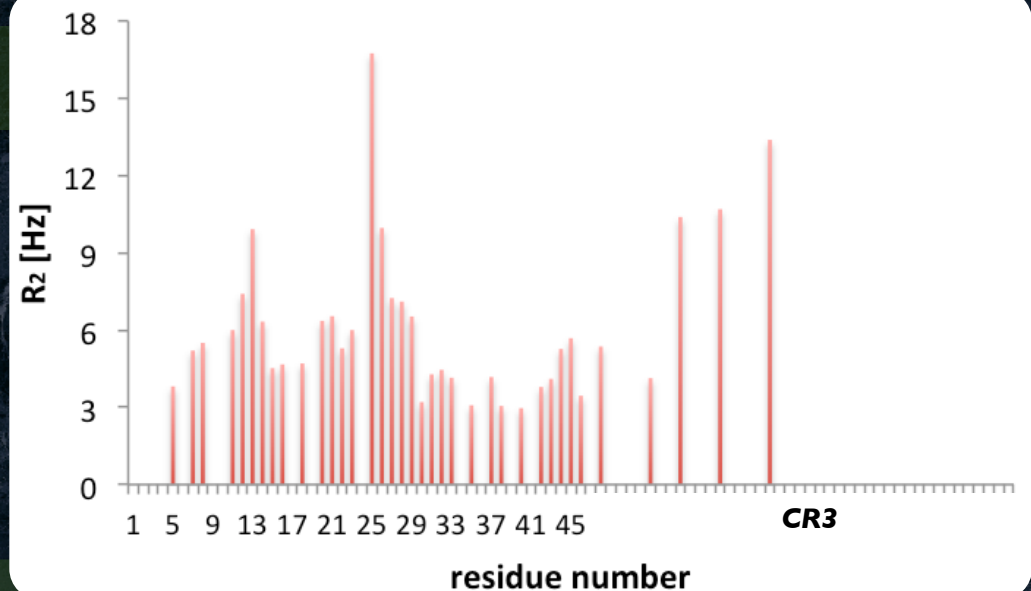
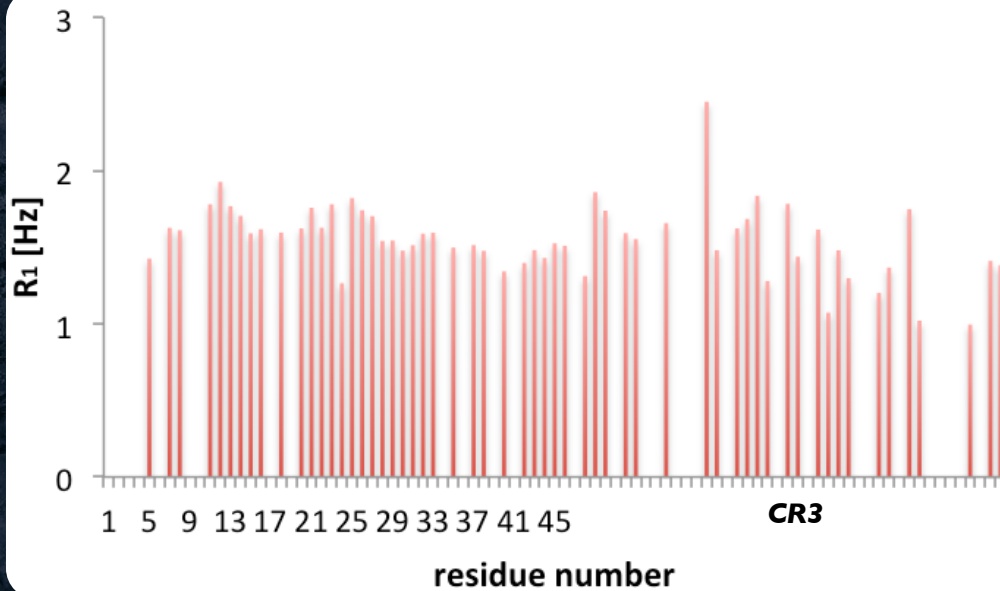
HPV 16 E7 oncoprotein



MHGDTH

CDSTLRLC
PICSQKP

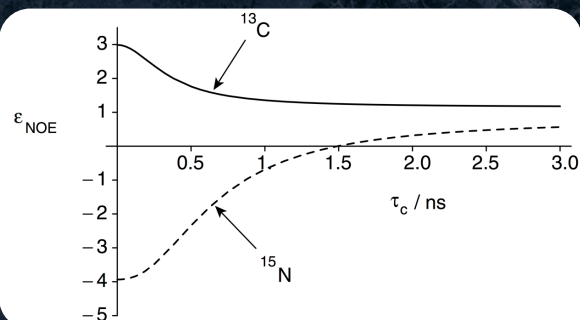
^{15}N relaxation experiments



QPETT

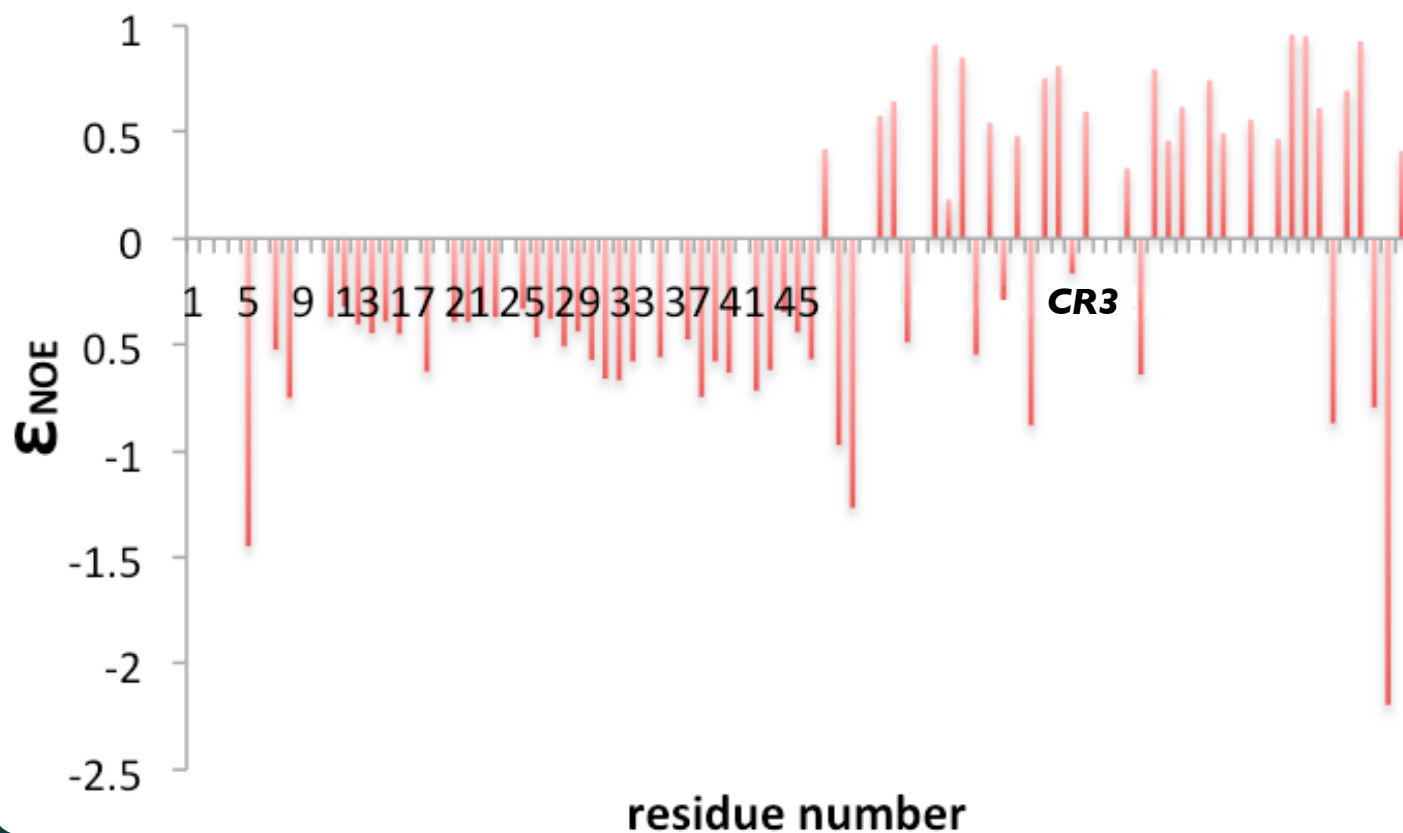
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VCPICSQKP

^1H - ^{15}N heteronuclear NOE



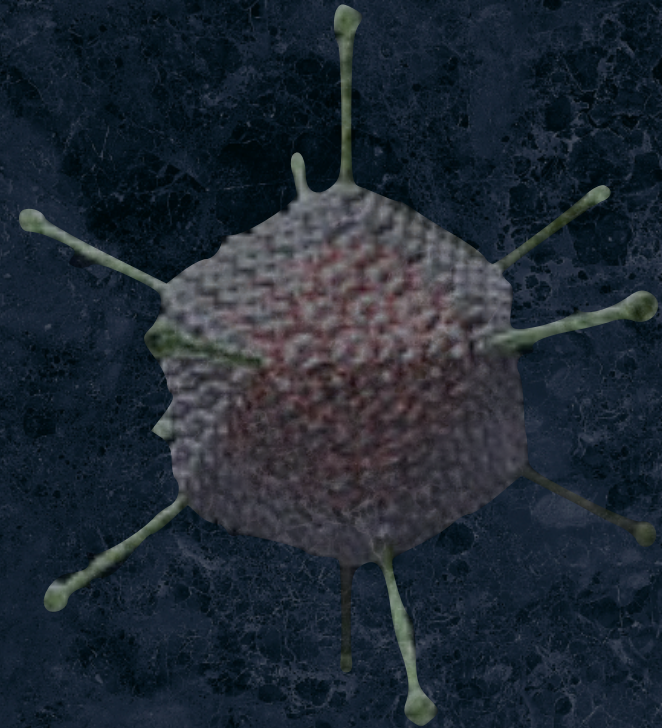
MHGDPTLHEY

QPETT



Adenovirus

- non-enveloped dsDNA virus
- involved in many infections:
 - acute respiratory
 - gastrointestinal
 - ocular
- can induce cancer, but also suppress tumorigenicity



Pelka, P. et al.(2008). Intrinsic Structural Disorder in Adenovirus E1A: a Viral Molecular Hub Linking Multiple Diverse Processes. Journal of Virology, 82(15), 7252–7263.

Adenovirus E1A protein



- can induce cancer (rodent)
- but also suppress tumorigenicity (human)
- around 50 known binding partners
- two main forms (289AA and 243 AA)



Pelka et al. (2008). Intrinsic Structural Disorder in Adenovirus E1A: a Viral Molecular Hub Linking Multiple Diverse Processes. Journal of Virology, 82(15), 7252–7263.

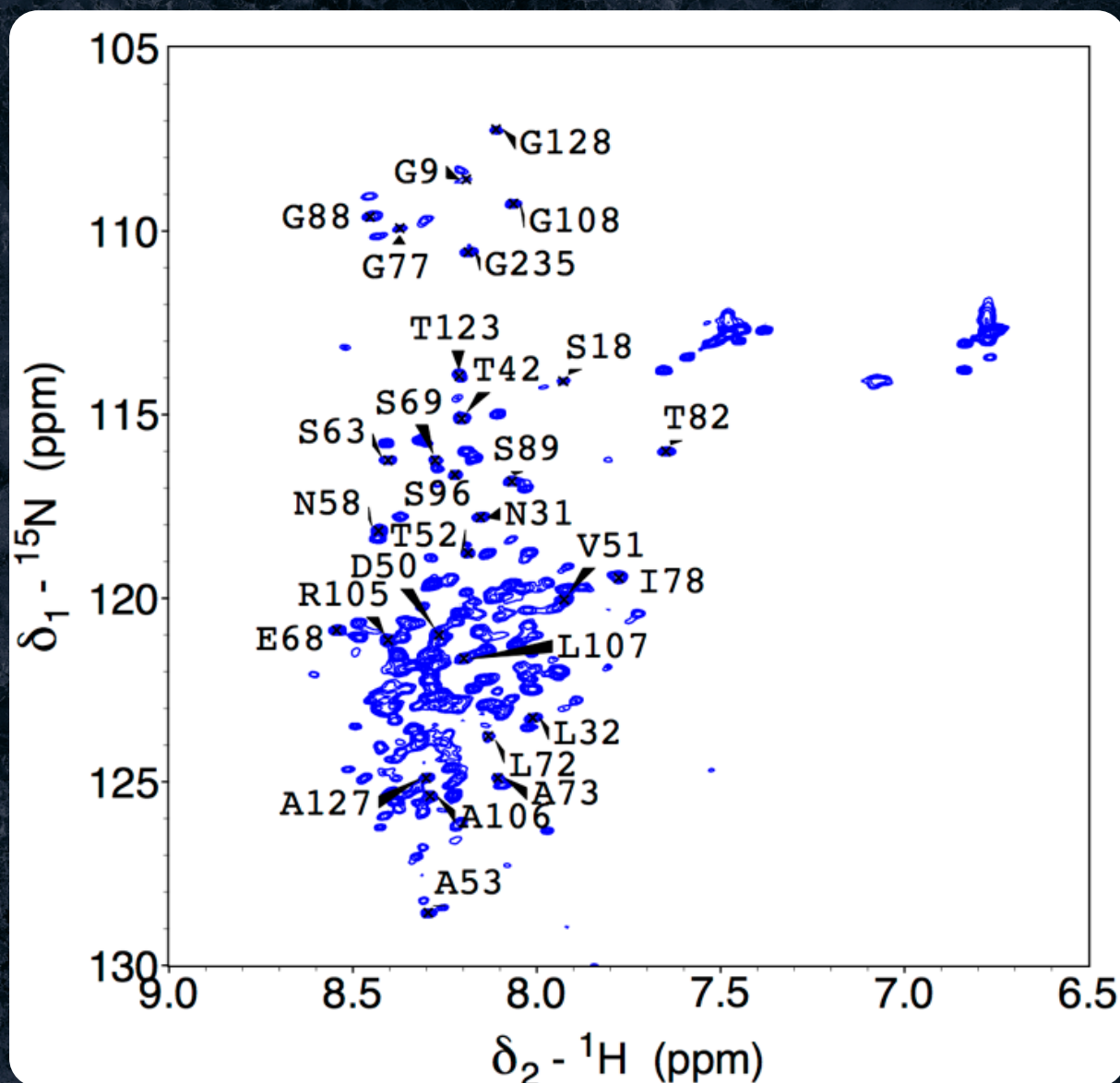
AdV 2/5 289R E1A protein

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GPVSMPNLVPEVIDLTCHEAGFP PSDDEDEEGEEFVLDYVGHPGHGCRSCHYH
RRNTGDPDIMCSLCYMRTCGMFVYSPVSEPEPEPEPEPEPARPTRR PKLVPAIL
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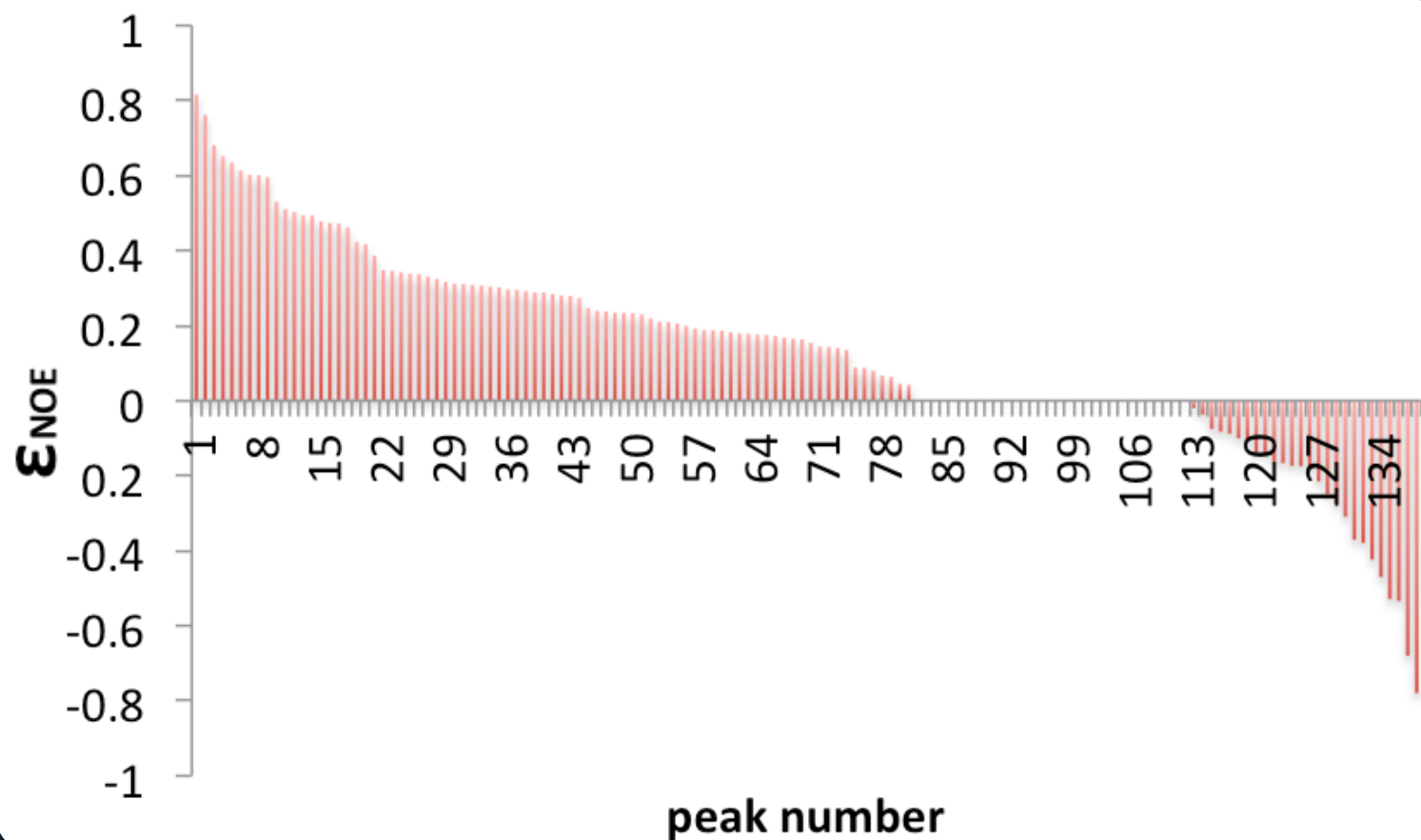
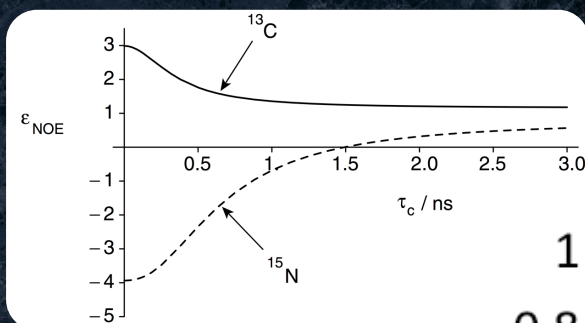
AdV 2/5 289R E1A protein

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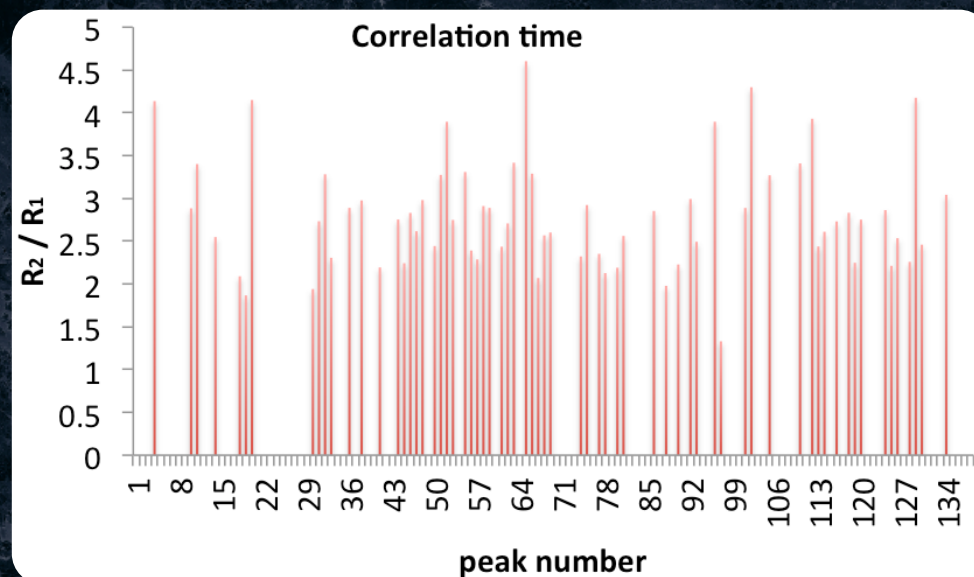
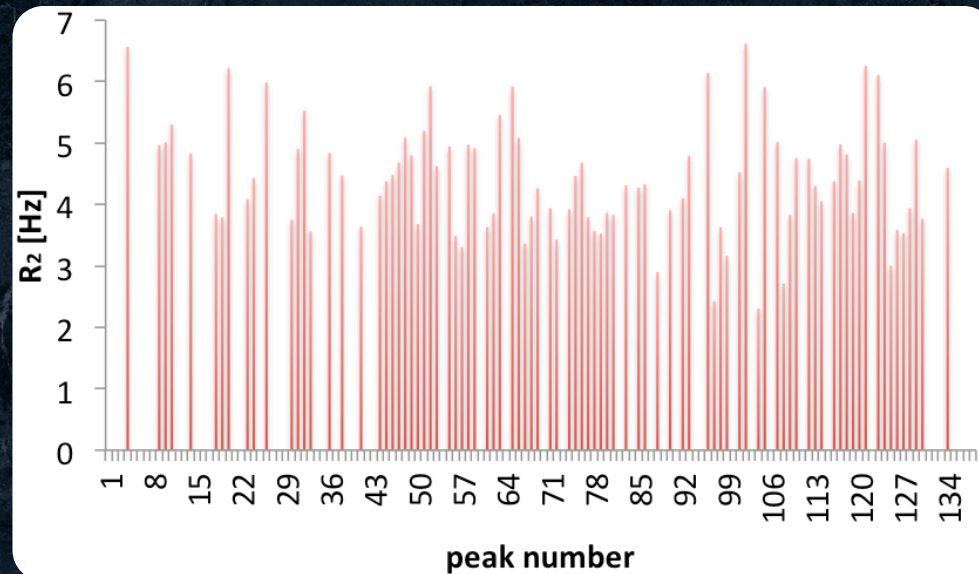
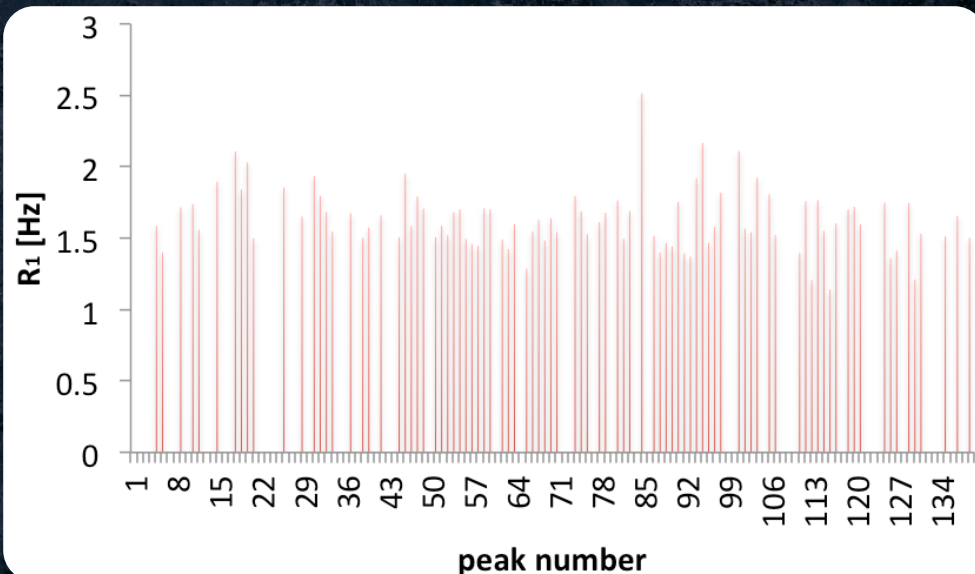
AdV 2/5 289R E1A protein



^1H - ^{15}N heteronuclear NOE



^{15}N relaxation experiments



Acknowledgement

- prof. Isabella C. Felli, prof. Roberta Pierattelli
- Eduardo Calçada
- CERM
- IDPbyNMR

Thank you!