

Supporting Information for

Interaction of the neurotransmitter, neuropeptide Y
(NPY), with phospholipid membranes:

Film balance and fluorescence microscopy studies

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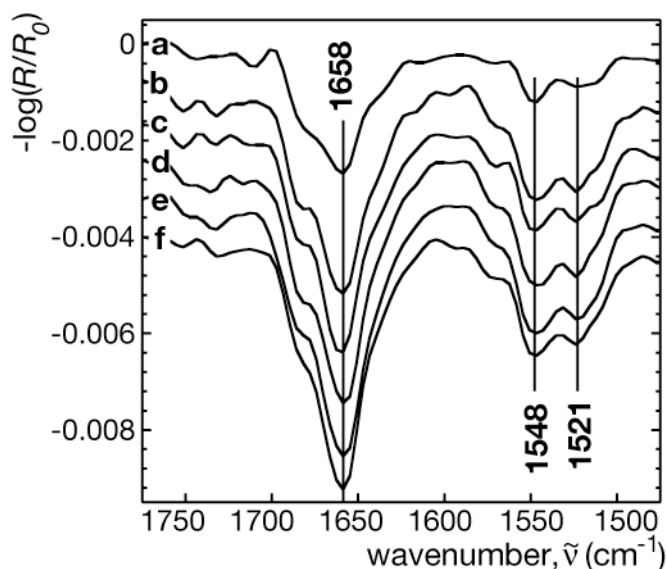


Figure S1: Amide region of external Fourier-transform infrared reflection-absorption spectra (FT-IRRA; p-polarization, incident angle, $\alpha_i = 40^\circ$, $T = 20^\circ\text{C}$) of a hNPY film on a water subphase at various states of compression (a: $\pi = 0$ mN/m, b: 8 mN/m, c: 12 mN/m and d: 16 mN/m). The FT-IRRA spectra show a strong, narrow absorption band at 1658 cm^{-1} , the amide I mode, which is mainly associated with the C=O stretching mode of the peptide backbone. The position of this band is conformationally sensitive¹ and indicates that hNPY adopts an α -helical secondary structure at the air/water interface. In addition, the amide II band at 1548 cm^{-1} also indicates an α -helical structure. The amide I band shows a high-frequency shoulder ($\sim 1685\text{ cm}^{-1}$) that may arise from a turn or a loop in the peptide.¹ Since hNPY has 5 tyrosine residues, the band in the amide II region at $\sim 1520\text{ cm}^{-1}$ can be attributed to C–H in-plane bending and C–C stretching vibrations of tyrosine.² Since there is no amide I absorption detected near 1630 cm^{-1} , it is rather unlikely that the band at 1520 cm^{-1} should be attributed to a β -sheet conformation.

The data have been collected in collaboration with Andreas Kerth and Alfred Blume at the Institute of Physical Chemistry at the Martin Luther University in Halle, Germany. A full account of this work will be presented elsewhere.³

REFERENCES

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