

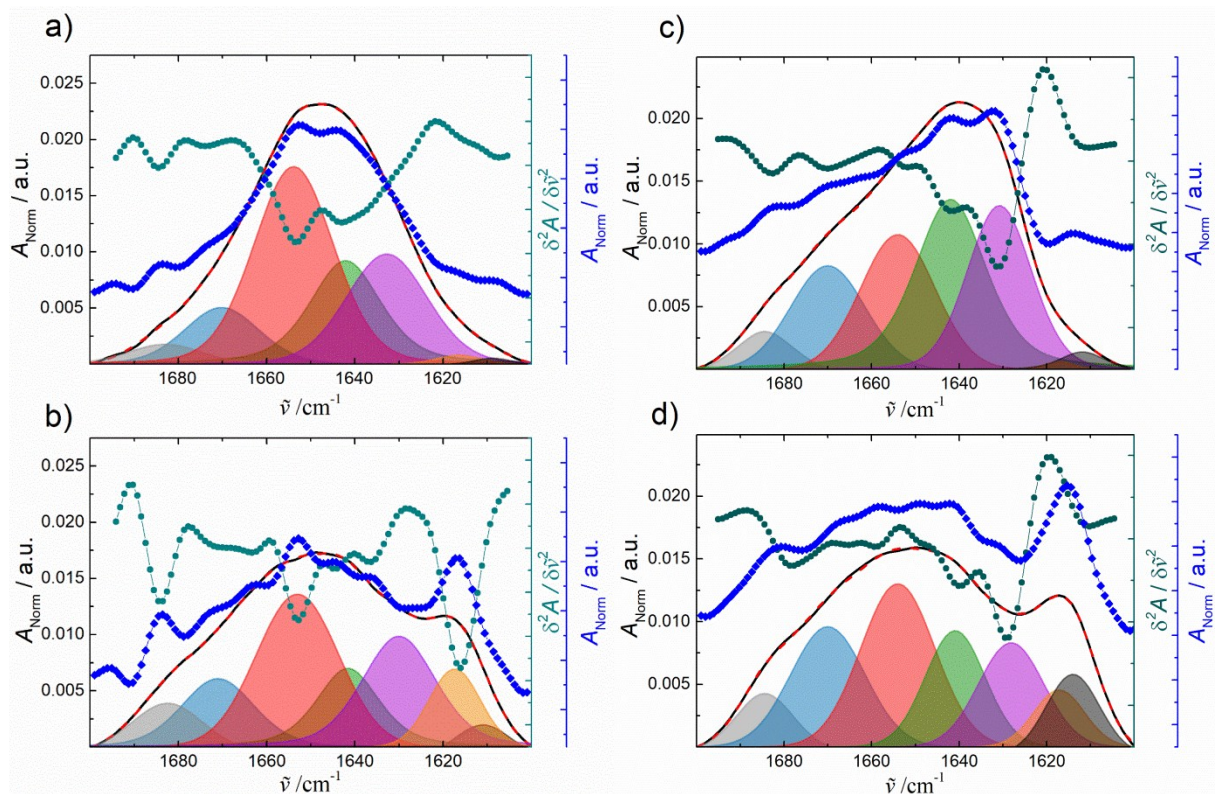
## Supporting Information

**Table SI 1.** Spectroscopically determined unfolding temperature and van't Hoff enthalpy change (\*) of GroEL, GroES and the GroEL-GroES complex (molar ratio of 1:2) in the absence and presence of different salts and cosolvents at ambient pressure. For GroES, two structural transitions (denoted 1, 2) could be observed.

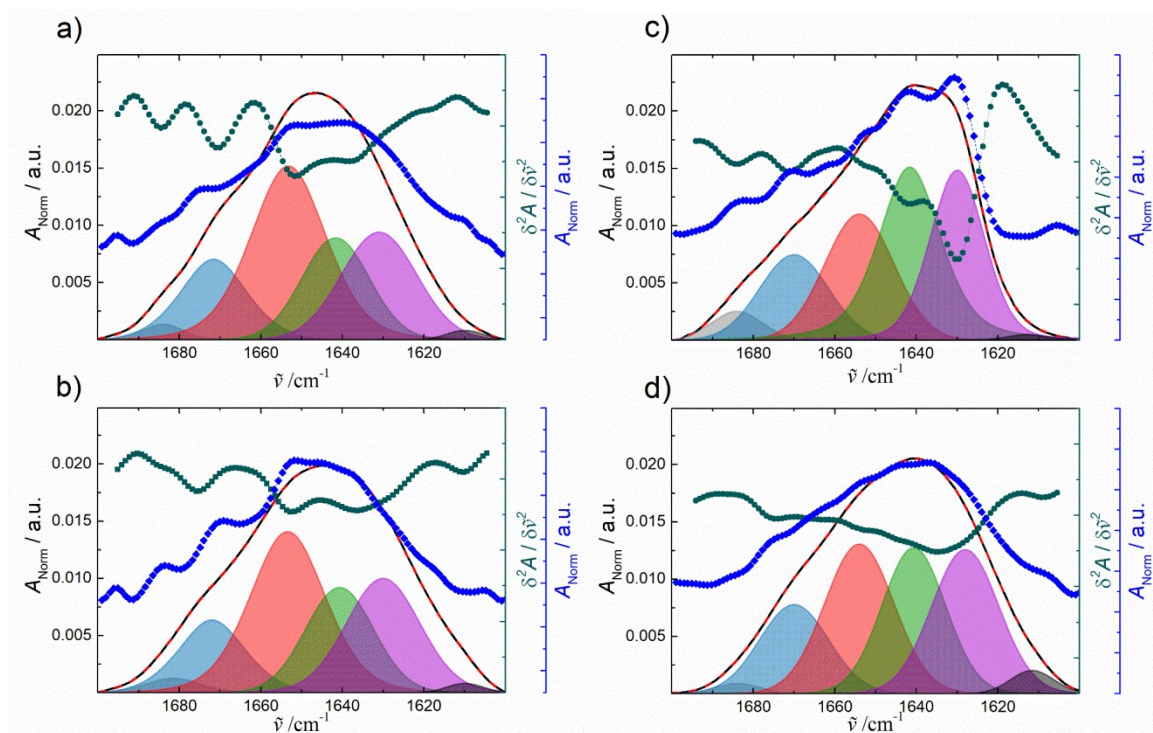
	$T_u / ^\circ\text{C}$	$\Delta H_{\text{vH,u}} / \text{kJ mol}^{-1}$
GroEL (buffer A)	$69.1 \pm 0.2$	$998 \pm 61$
GroEL (buffer A + 3 mM ADP)	$67.8 \pm 0.2$	$732 \pm 89$
GroEL (buffer B)	$72.0 \pm 0.6$	$821 \pm 84$
GroEL (buffer B + 3 mM ATP)	$70.9 \pm 0.5$	$788 \pm 79$
GroEL (buffer B + 3 mM ATP + 10 mM $\text{Mg}^{2+}$ )	$70.6 \pm 0.4$	$835 \pm 37$
GroEL (buffer B + 10 mM $\text{Mg}^{2+}$ )	$71.8 \pm 0.6$	$916 \pm 77$
GroEL (buffer B + 1 M $^{13}\text{C}$ -urea)	$65.1 \pm 0.3$	$917 \pm 170$
GroEL (buffer B + 1 M TMAO)	$73.7 \pm 0.6$	$732 \pm 91$
GroES (buffer B)	1) $82.1 \pm 0.3$ 2) $87.1 \pm 0.6$	1) $691 \pm 92$ 2) $669 \pm 286$
GroES (buffer B + 1 M $^{13}\text{C}$ -urea)	1) $80.0 \pm 1.2$ 2) $83.5 \pm 1.0$	1) $829 \pm 141$ 2) $1028 \pm 245$
GroES (buffer B + 1 M TMAO)	1) $84.8 \pm 1.2$ 2) $87.8 \pm 0.6$	1) $822 \pm 133$ 2) $671 \pm 132$
GroEL-GroES (buffer B)	$71.4 \pm 0.5$	$1007 \pm 144$
GroEL-GroES (buffer B + 3 mM ATP + 10 mM $\text{Mg}^{2+}$ + 1 M $^{13}\text{C}$ -urea)	$67.2 \pm 0.4$	$863 \pm 70$
GroEL-GroES (buffer B + 3 mM ATP + 10 mM $\text{Mg}^{2+}$ + 1 M TMAO)	$75.9 \pm 0.5$	$879 \pm 61$

**Table SI 2.** Spectroscopically determined unfolding pressures,  $p_u$ , and volume change of (partial) unfolding,  $\Delta V_u$ , of GroES in the absence and presence of different cosolvents at 25 °C.

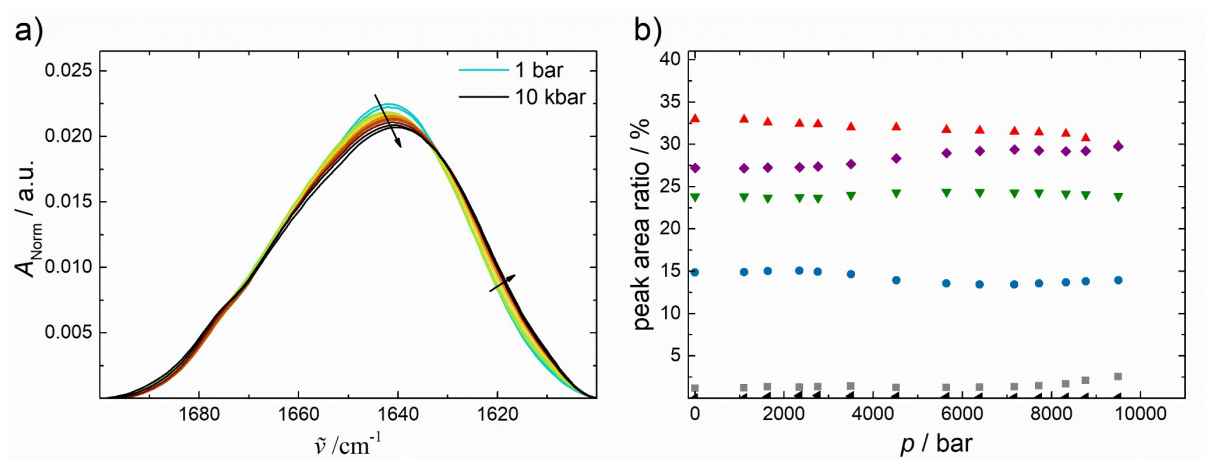
	$p_u / \text{bar}$	$\Delta V_u / \text{cm}^3 \text{mol}^{-1}$
GroES (buffer B)	$5395 \pm 298$	$-52.9 \pm 11.8$
GroES (buffer B + 1 M TMAO)	$5413 \pm 321$	$-54.8 \pm 7.8$
GroES (buffer B + 1 M $^{13}\text{C}$ -urea)	$3552 \pm 85$	$-112.28 \pm 14.6$



**Figure SI 1.** Curve fitting results of the temperature-dependent FTIR data of GroEL and GroES in neat buffer at ambient pressure: Amide I' band region (black line), the corresponding fit (red dashed line), FSD (blue dotted (square) line) and 2<sup>nd</sup> derivative spectra (dark cyan dotted line) of GroEL for 25 °C (a) and 86 °C (b) at ambient pressure and of GroES at 25 °C (c) and 96 °C (d) at ambient pressure: intermolecular  $\beta$ -sheets (grey), turns and loops (blue),  $\alpha$ -helices (red), random coils (green), intramolecular  $\beta$ -sheets (purple), intermolecular  $\beta$ -sheets (orange) and side chains (black).



**Figure SI 2.** Curve fitting results of the pressure-dependent FTIR data of GroEL and GroES in neat buffer at 25 °C. Amide I' band region (black line), the corresponding fit (red dashed line), FSD (blue dotted (square) line) and 2<sup>nd</sup> derivative spectra (dark cyan dotted line) of GroEL for 1 bar (a) and 10 kbar (b) at 25 °C and of GroES for 1 bar (c) and 10 kbar (d) at 25 °C: intermolecular  $\beta$ -sheets (grey); turns and loops (blue);  $\alpha$ -helices (red); random coils (green); intramolecular  $\beta$ -sheets (purple) and side chains (black).



**Figure SI 3.** Pressure-dependent FTIR absorption data of the GroEL-GroES complex in neat buffer (buffer B + 3 mM ATP + 10 mM  $\text{Mg}^{2+}$ ) at 25 °C. a) Pressure-dependent changes of the normalized amide I' band region of the GroEL-GroES complex and b) corresponding changes in secondary structure elements by pressure: intermolecular  $\beta$ -sheets (1683  $\text{cm}^{-1}$ , grey); turns and loops (1672  $\text{cm}^{-1}$ , blue);  $\alpha$ -helices (1653  $\text{cm}^{-1}$ , red); random coils (1640  $\text{cm}^{-1}$ , green); intramolecular  $\beta$ -sheets (1632  $\text{cm}^{-1}$ , purple) and side chains (1610  $\text{cm}^{-1}$ , black).