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Motivation

Multidimensional fifth order spectroscopy (R^5 -2D-ES) is a sophisticated and specialized tool for probing bi-exciton dynamics [1,2]. Here, we present an analysis of fifth order signals emitted into $\pm 2K_1 \mp 2K_2 + K_3$ phase matching directions. In an effort to model fifth order electronic spectra of tractably small systems, we develop a dynamical model for molecular dimers and trimers. **We show that fifth order two-dimensional spectroscopy allows to track and pinpoint population transfer- and annihilation dynamics of bi-excitons, without the need to perform and analyse intensity dependent experiments.**

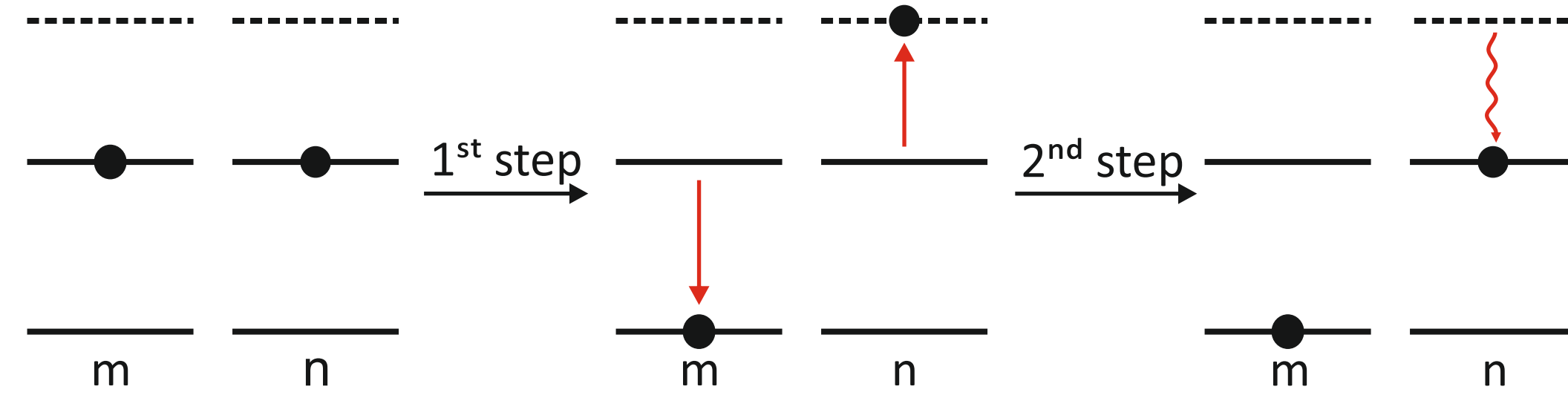


Fig. 1: Microscopic picture of EEA [3]: (1) excitation energy is resonantly transferred between two adjacent sites m and n . This transfer stimulates one site back to the ground state whereas the other is promoted to a higher lying state. (2) Phonon-assisted non-radiative relaxation back to the first excited state.

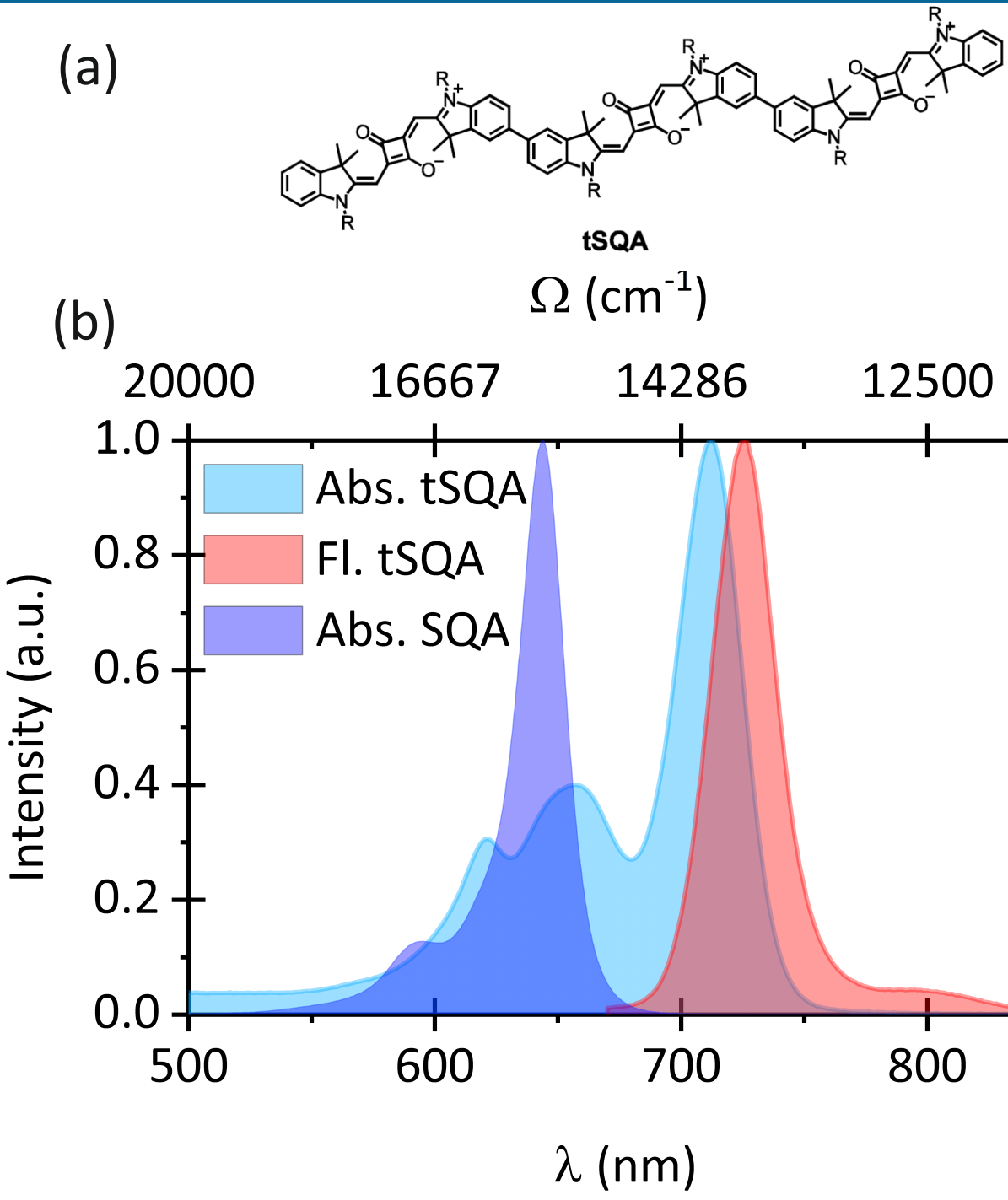


Fig. 2:
(a) Structure of tSQA
(b) Linear spectra of SQA and tSQA

Multidimensional line shapes of the fifth order response

Fifth order signals:

$$R_{NR}^{(5)}(t_1, t_2, t_3) = (i)^5 \langle \langle \mu | \check{G}(t_3) \check{\mu}^{(+)} \check{G}(t_2) \check{\mu}^{(-)} \check{\mu}^{(-)} \check{G}(t_1) \check{\mu}^{(+)} \check{\mu}^{(+)} | gg \rangle \rangle$$

$$R_R^{(5)}(t_1, t_2, t_3) = (i)^5 \langle \langle \mu | \check{G}(t_3) \check{\mu}^{(+)} \check{G}(t_2) \check{\mu}^{(+)} \check{\mu}^{(+)} \check{G}(t_1) \check{\mu}^{(-)} \check{\mu}^{(-)} | gg \rangle \rangle$$

Dipole superoperator actions:

Green's function:

$$\check{\mu}^{(+)} \hat{X} = \sum_{n=1}^N d_n (\hat{B}_n^\dagger \hat{X} - \hat{X} \hat{B}_n^\dagger)$$

$$\check{G}(t) = \exp\left(\left(\check{D} + \check{A}\right)t\right)$$

$$\check{\mu}^{(-)} \hat{X} = \sum_{n=1}^N d_n (\hat{B}_n \hat{X} - \hat{X} \hat{B}_n)$$

\check{D} ... generator for transport: Redfield and Förster

\check{A} ... generator for annihilation dynamics

d_n ... transition dipole moment of molecule n

\hat{B}_n^\dagger (\hat{B}_n) ... creation (annihilation) operator of excitation at site n

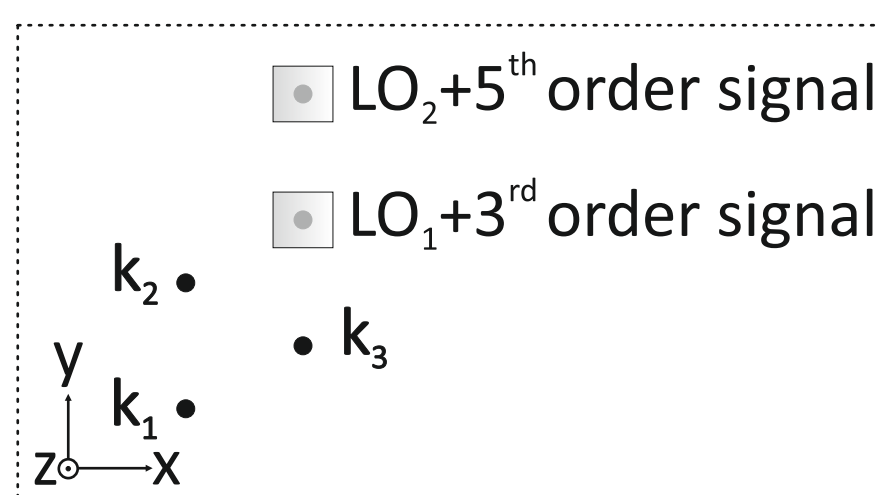


Fig. 4: Schematic of the experimental phase matching geometry

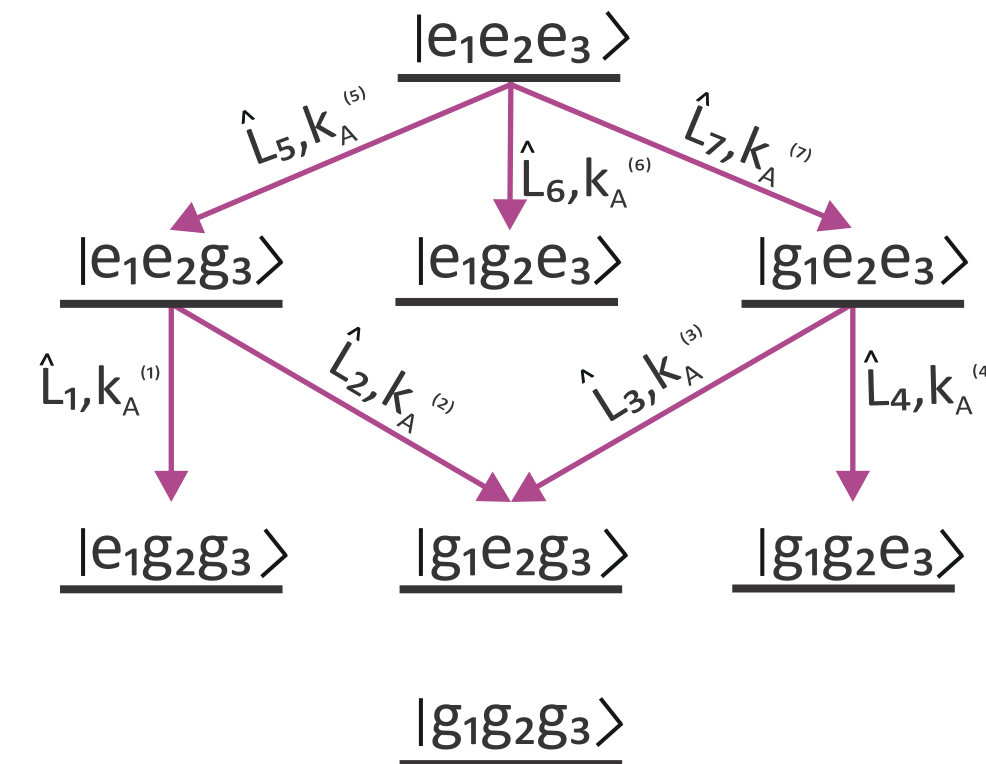


Fig. 3: EEA in a molecular trimer is modeled as unidirectional transport to lower excitonic manifolds induced by operators $\{L_i\}$ and rates $k_A^{(i)}$, as indicated by purple arrows.

$2k_1 - 2k_2 + k_3$:	GSB	ESA		SE		SE	ESA
t_2 dynamics	/	Coherence $i \neq m$	Transport $m \rightarrow n$	Coherence $i \neq j$	Transport $l \rightarrow m$	Exciton	Exciton Annihilation
bi-exciton dynamics							
single-exciton dynamics						Energy level diagram	

Fig. 2: Liouville space pathways (LSP) relevant for the calculation of the non-rephasing $K_{NR} = +2K_1 - 2K_2 + K_3$ signal of molecular dimers and trimers. Single-, bi- and tri-exciton states are labeled by $\{|a_i\rangle\}$, $\{|\alpha_i\rangle\}$ and $\{|f_i\rangle\}$, respectively. The energy level diagram of the considered molecular trimer is depicted at the bottom right. GSB: Ground state bleach, ESA: excited state absorption, SE: stimulated emission.

Molecular dimer:

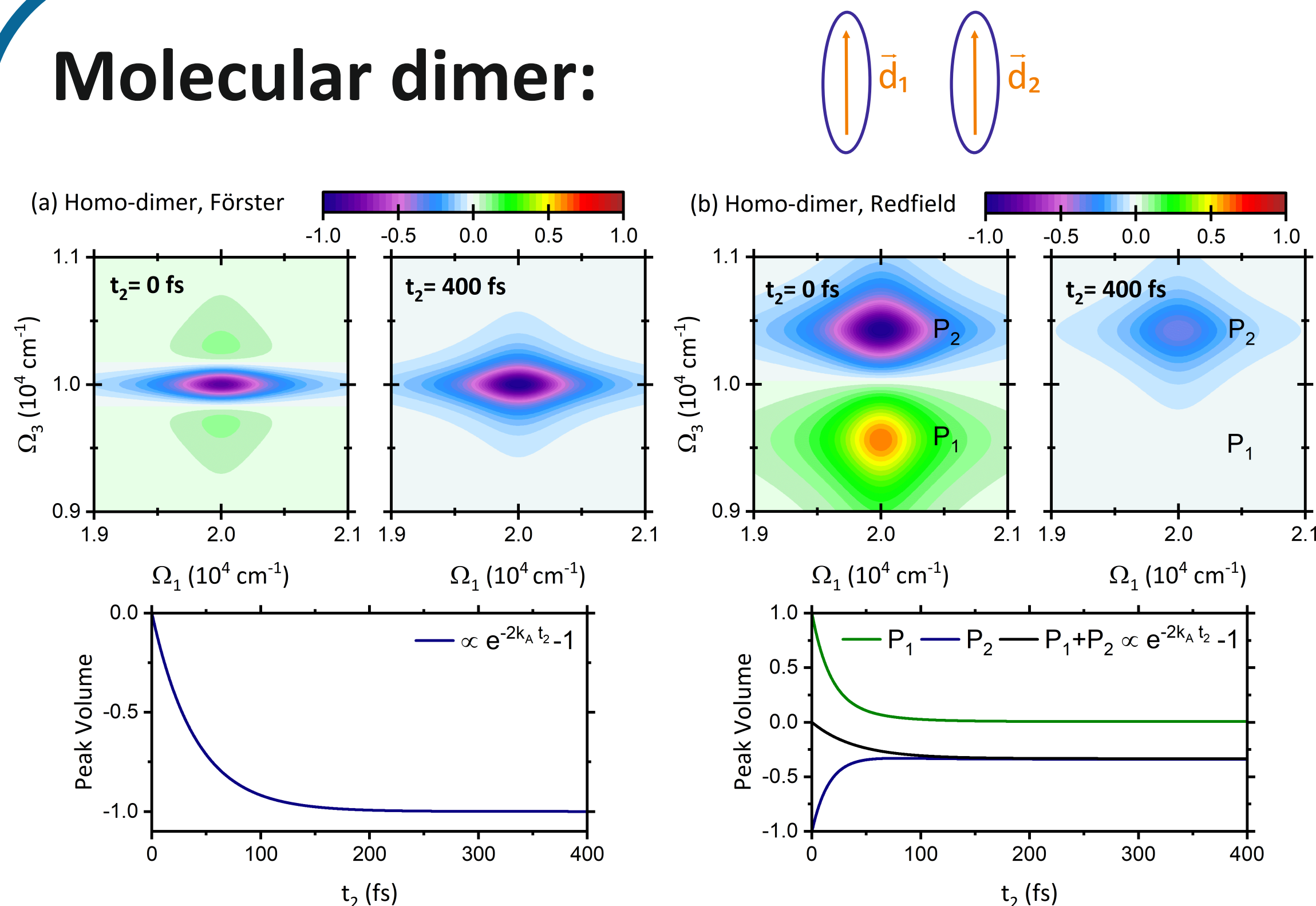


Fig. 5: R^5 -2D-ES signal (top panels) of a weakly (a) and strongly (b) coupled homo-dimer for two different waiting times. The bottom panel in both (a) and (b) shows the evolution of the peak volume $V(t_2)$ versus the waiting time t_2 .
Parameter:
 $E = 10^4 \text{ cm}^{-1}$, $k_A^{-1} = 80 \text{ fs}$; Förster: $k_{2 \rightarrow 1}^{-1} = k_{1 \rightarrow 2}^{-1} = 400 \text{ fs}$; Redfield: $k_{a_2 \rightarrow a_1}^{-1} = 17 \text{ fs}$, $k_{a_1 \rightarrow a_2}^{-1} = 980 \text{ fs}$, $J = 420 \text{ cm}^{-1}$.

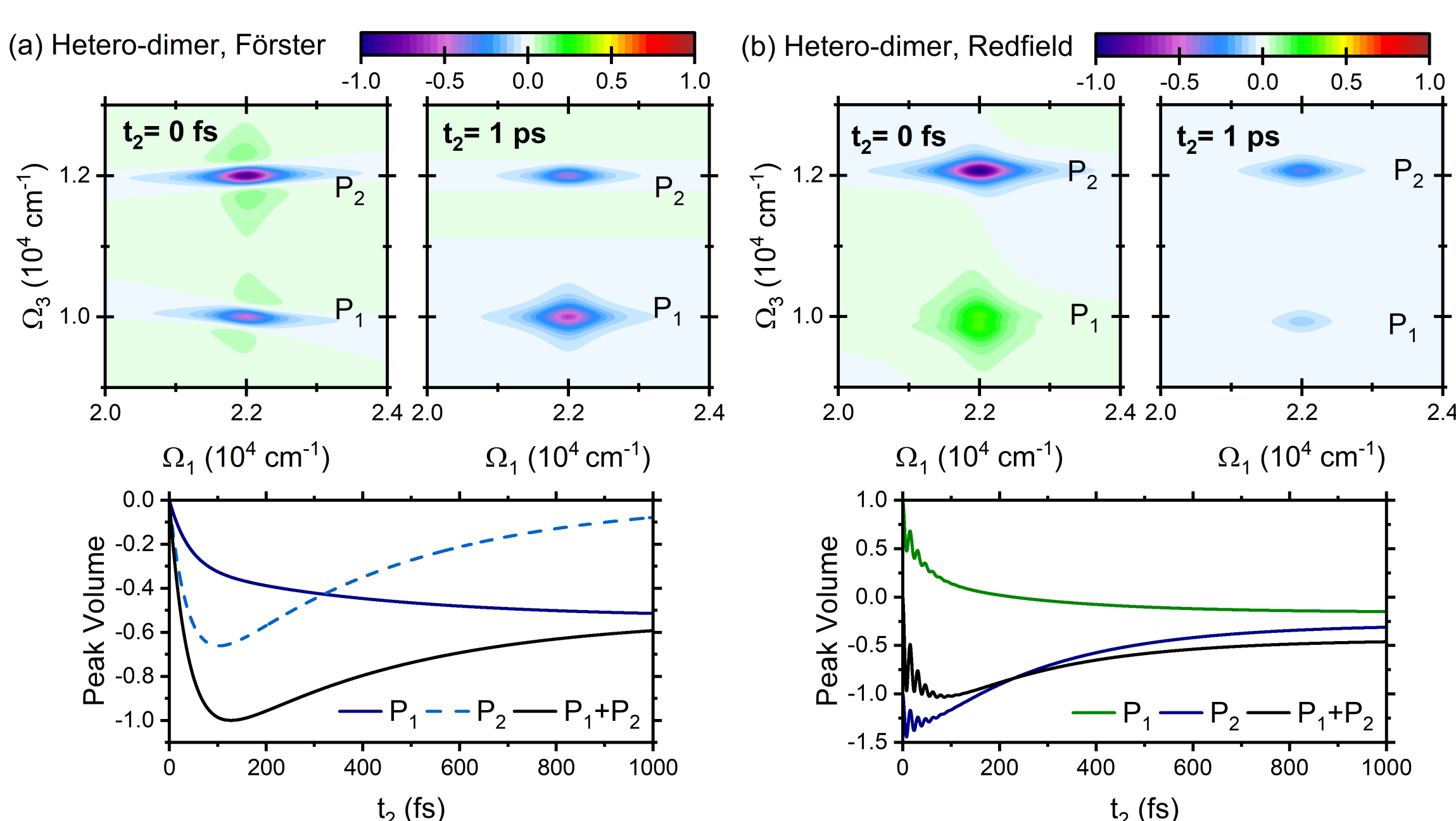


Fig. 6: R^5 -2D-ES signal (top panels) of a weakly (a) and strongly (b) coupled hetero-dimer for two different waiting times. The bottom panel in both (a) and (b) shows the evolution of the peak volume versus the waiting time t_2 .
Parameter:
 $E_1 = 10^4 \text{ cm}^{-1}$, $E_2 = 1.2 \cdot 10^4 \text{ cm}^{-1}$, $d = 2d$, $k_A^{-1} = 80 \text{ fs}$; Förster: $k_{2 \rightarrow 1}^{-1} = 400 \text{ fs}$, $k_{1 \rightarrow 2}^{-1} = 0 \text{ fs}$; Redfield: $k_{a_2 \rightarrow a_1}^{-1} = 265 \text{ fs}$, $k_{a_1 \rightarrow a_2}^{-1} = 0 \text{ fs}$, $J = 374 \text{ cm}^{-1}$.

Molecular trimer:

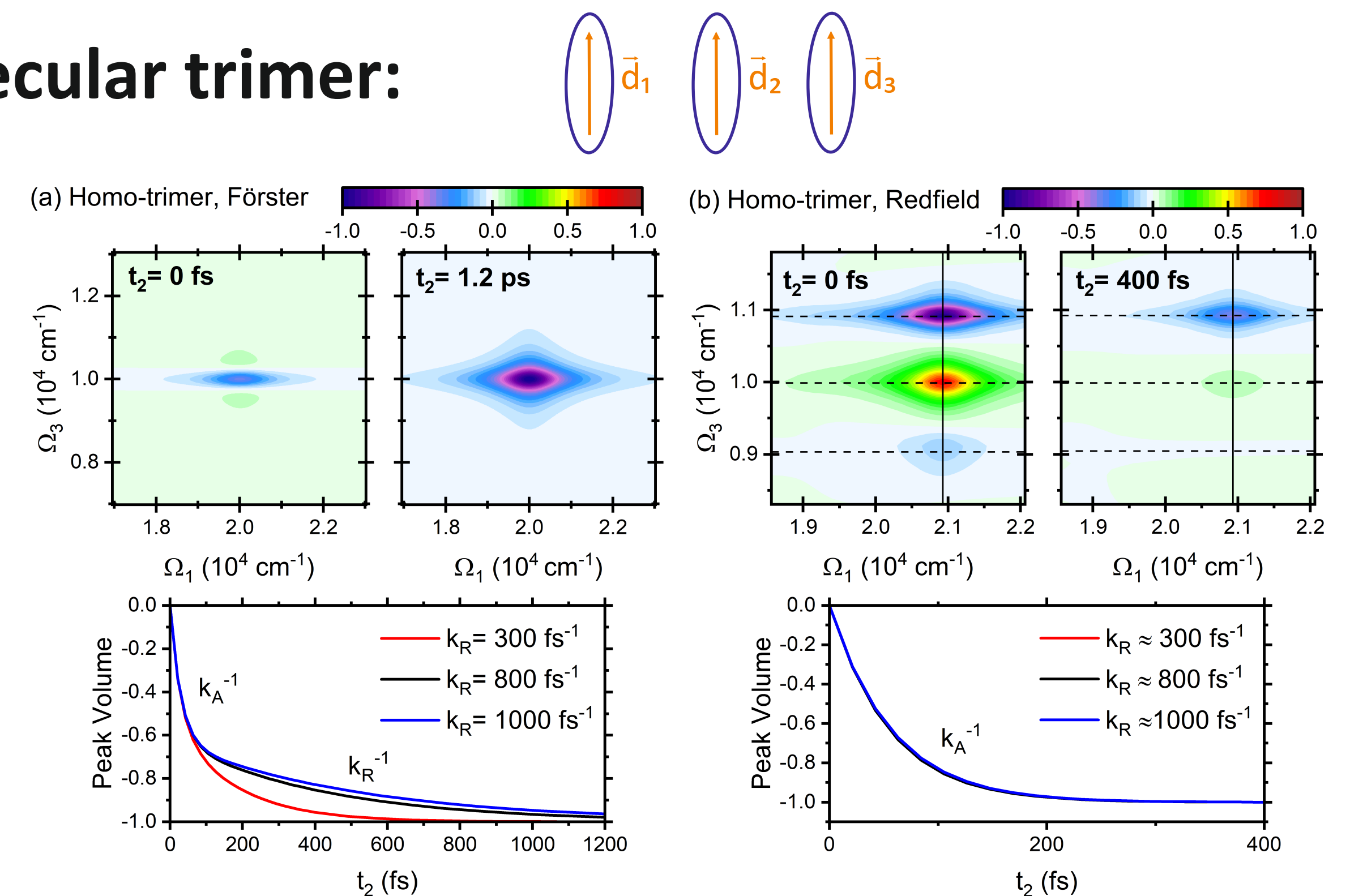


Fig. 7: R^5 -2D-ES signal of a weakly (top panels of a) and strongly (top panels of b) coupled homo-trimer for two different waiting times. The bottom panel in both (a) and (b) shows the evolution of the peak volume for different relaxation rates versus the waiting time t_2 .
Parameter:
 $E = 10^4 \text{ cm}^{-1}$, $k_A^{-1} = 60 \text{ fs}$; Förster: $k^{-1} = 300 \text{ fs}$; Redfield: $k_{a_2 \rightarrow a_1}^{-1} = k_{a_3 \rightarrow a_2}^{-1} = 300 \text{ fs}$, $k_{a_1 \rightarrow a_2}^{-1} = k_{a_2 \rightarrow a_3}^{-1} = k_{a_1 \rightarrow a_3}^{-1} = 0 \text{ fs}$, $J_{12} = J_{23} = 630 \text{ cm}^{-1}$, $J = 80 \text{ cm}^{-1}$.

Conclusion:

- The signal from a homo-dimer is directly related to the annihilation rate.
- The peak volume evolution of the hetero-dimer shows a more involved dependency on both annihilation and transport rates.
- The timescale of exciton diffusion affects the peak volume decay of a weakly coupled trimer, as excitons must migrate to adjacent sites to annihilate. The highly delocalized excitons of a strongly coupled trimer, however, do not allow for more than one stable exciton within the delocalization length and thus no exciton diffusion timescale is visible in the peak volume decay.
- Annihilation dynamics is essential for the appearance of an integrated signal of homo-aggregates of arbitrary length [1].

$$V(t_2) = 8d_1^2 d_2^2 \left[\left(d_1^2 + d_2^2 \right) \left(G_{\alpha\alpha, \alpha\alpha}(t_2) - G_{gg, gg}(t_2) \right) + \left(d_1^2 - d_2^2 \right) \left(G_{22, 22}(t_2) - G_{11, 11}(t_2) - G_{11, 22}(t_2) + G_{22, 11}(t_2) - G_{22, \alpha\alpha}(t_2) + G_{11, \alpha\alpha}(t_2) \right) \right]$$

Analytical expression for the peak volume dynamics of a weakly coupled hetero dimer. The signal is directly related to the annihilation rate, in the limit of $d_1^2 = d_2^2$.