

Can an entirely negative fluorine in a molecule, viz. perfluorobenzene, interact attractively with the entirely negative site(s) on another molecule(s)? Like liking like!

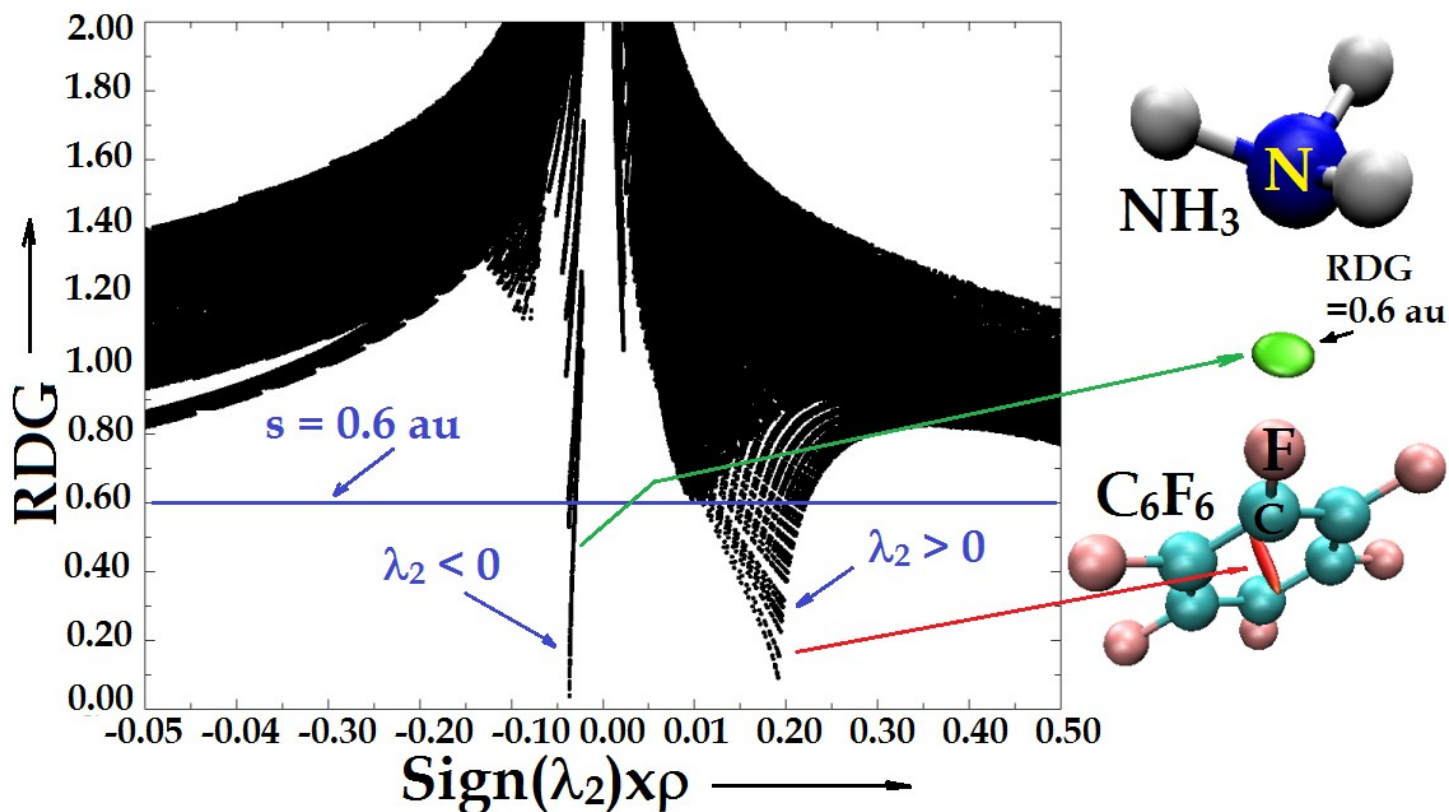
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Supplementary Information

Fig. S1: RDG vs. $\text{Sign}(\lambda_2)\times\rho$ (left) and RDG = 0.6 a.u. isosurface (right) NCI graphical plots for $\text{H}_3\text{N}\cdots\text{FC}_6\text{F}_5$. The spikes in the $\lambda_2 < 0$ and $\lambda_2 > 0$ regions represent to the bonded (attractive) and non-bonded (repulsive) interactions, respectively. The respective $s = 0.6$ a.u. RDG isosurfaces painted in green and red are depicted in Fig. 2 (see the ball and stick models). (Also, see Fig. 1 for conformational details, and that the dimer labeling in Fig. 1 is in consistent with that shown in Fig. 2 for QTAIM and RDG based molecular and RDG plots, respectively).



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Fig. S2: RDG vs. $\text{Sign}(\lambda_2)\times\rho$ plot for $\text{H}_2\text{O}\cdots\text{FC}_6\text{F}_5$ a), $\text{H}_2\text{O}\cdots\text{FC}_6\text{F}_5$ b), $\text{H}_3\text{CF}\cdots\text{FC}_6\text{F}_5$ c), $\text{H}_3\text{CF}\cdots\text{FC}_6\text{F}_5$ d), $\text{HF}\cdots\text{FC}_6\text{F}_5$ e) and $\text{HF}\cdots\text{FC}_6\text{F}_5$ f) (values in a.u.). The spikes in the $\lambda_2 < 0$ and $\lambda_2 > 0$ regions represent to the bonded (attractive) and non-bonded (repulsive) interactions, respectively. The respective $s = 0.6$ a.u. RDG isosurfaces painted in green and red are depicted in Fig. 2 (see the ball and stick models). (Also, see Fig. 1 for conformational details, and that the dimer labeling in Fig. 1 is in consistent with that shown in Fig. 2 for QTAIM and RDG based molecular and RDG plots, respectively).

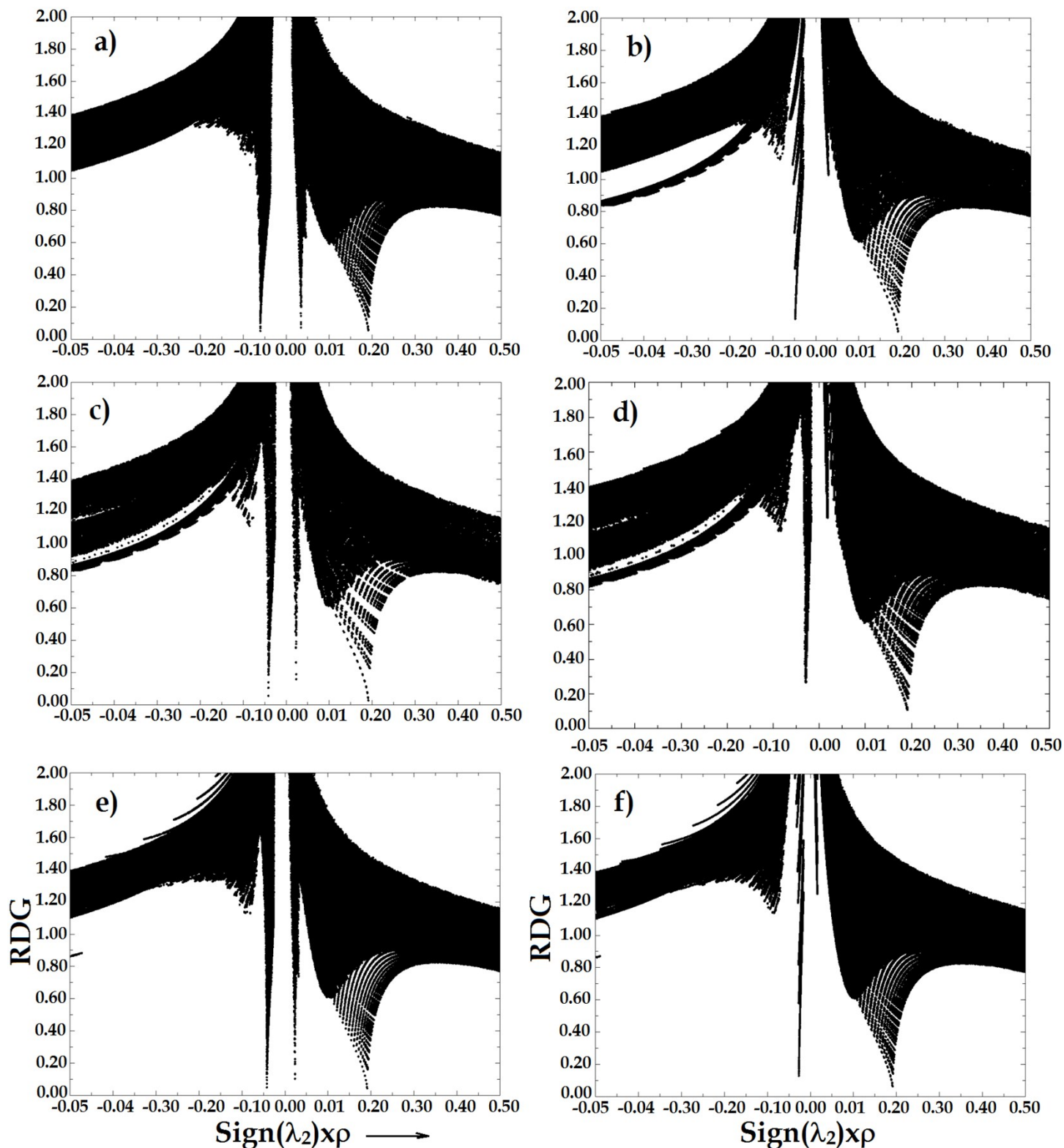


Fig. S3: RDG vs. $\text{Sign}(\lambda_2) \times \rho$ plot for $\text{H}_2\text{CO} \cdots \text{FC}_6\text{F}_5$ (h), $\text{H}_2\text{CO} \cdots \text{FC}_6\text{F}_5$ (i), $\text{C}_5\text{F}_5\text{N} \cdots \text{FC}_6\text{F}_5$ (j), $\text{C}_4\text{F}_4\text{N}_2 \cdots \text{FC}_6\text{F}_5$ (k), $\text{C}_4\text{F}_4\text{N}_2 \cdots \text{FC}_6\text{F}_5$ (l), and $(\text{CF})_4\text{N}_2 \cdots \text{FC}_6\text{F}_5$ (m) (values in a.u.). The spikes in the $\lambda_2 < 0$ and $\lambda_2 > 0$ regions represent to the bonded (attractive) and non-bonded (repulsive) interactions, respectively. The respective $s = 0.6$ a.u. RDG isosurfaces painted in green and red are depicted in Fig. 2 (see the ball and stick models). (Also, see Fig. 1 for conformational details, and that the dimer labeling in Fig. 1 is in consistent with that shown in Fig. 2 for QTAIM and RDG based molecular and RDG plots, respectively).

