

Fluorine Conformational Effects Characterized by Energy Decomposition Analysis

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ELECTRONIC SUPPLEMENTARY INFORMATION

Table S1. IQA energy components at the HF-D3/cc-pVTZ (sum for all the atoms and interatomic interactions) for the energy difference (kcal/mol) between the two conformers (gauche/anti or trans/cis) in 1,2-difluoroethane and related small systems.

$\Delta E = E_{gauche} - E_{anti}$	ΔE_{net}	$\Delta E_{int,disp}$	$\Delta E_{int,xc}$	$\Delta E_{int,class}$	ΔE_{IQA}
1: 1,2-difluoroethane	3.7	-0.1	-1.9	-2.3	-0.5
2: 2-fluoroethyl-acetate	0.2	-0.1	-2.2	1.6	-0.5
3: 3-fluoropropanal	-10.9	-0.0	0.9	8.9	-1.2
4: <i>N</i> -(2-fluoroethyl)acetamide	-24.6	-0.3	-2.3	25.0	-2.1
5: 2-(2-fluoroethyl)isoindoline-1,3-dione	-0.3	-0.3	-2.2	2.3	-0.5
6: 2-fluoroethan-1-aminium	-24.8	-0.6	1.9	16.2	-7.3
7: 1-(2-fluoroethyl)pyridin-1-ium	-20.4	-0.4	-0.8	16.5	-5.1
$\Delta E = E_{trans} - E_{cis}$	ΔE_{net}	$\Delta E_{int,disp}$	$\Delta E_{int,xc}$	$\Delta E_{int,class}$	ΔE_{IQA}
8: 1-fluoropropan-2-one	-30.8	-0.0	5.0	23.1	-2.7
9: methyl-2-fluoroacetate	2.1	-0.0	1.4	-3.5	0.0
10: 2-fluoro- <i>N</i> -methylacetamide	-4.8	-0.2	8.7	-9.5	-5.8

Table S2. Largest differences in the atomic energies (kcal/mol) between the gauche and anti $F_2-C_1-C_5-X$ conformers in CH_2F-CH_2X systems ($\Delta E = E_{gauche} - E_{anti}$). Ball-and-stick models of the gauche conformer including atom numbering.

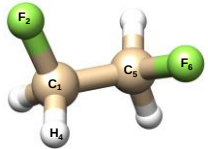
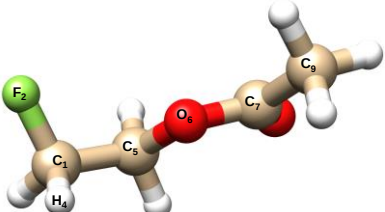
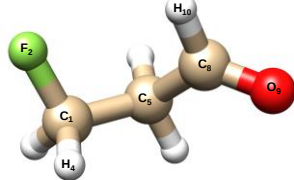
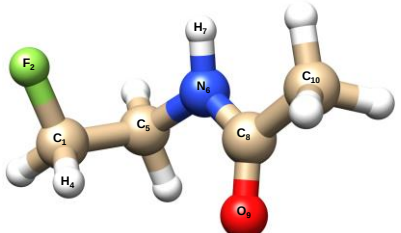
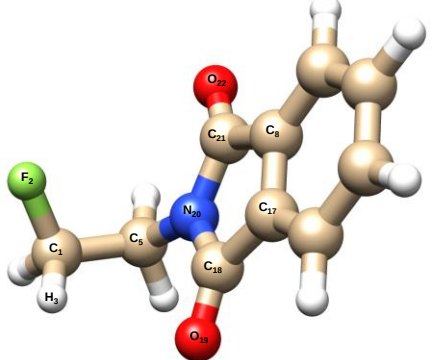
$\Delta E = E_{gauche} - E_{anti}$	ΔE_{net}		$\Delta E_{int,clas}^{AB}$		$\Delta E_{int,xc}^{AB}$	
1: 1,2-difluoroethane 	F ₂	1.1	F ₂ ...F ₆	12.1	C ₁ ...C ₅	-3.2
	F ₆	1.0	C ₁ ...F ₆	-8.0	C ₁ ...F ₆	1.0
	C ₁	1.0	F ₂ ...C ₅	-7.9	F ₂ ...C ₅	1.0
			C ₁ ...C ₅	6.9	F ₂ ...F ₆	-0.9
					C ₁ ...H ₄	0.8
					C ₅ ...H ₈	0.8
2: 2-fluoroethyl-acetate 	O ₆	1.3	F ₂ ...O ₆	20.8	C ₁ ...C ₅	-3.0
			F ₂ ...C ₇	-14.0	F ₂ ...O ₆	-1.6
			C ₁ ...O ₆	-11.9	C ₁ ...O ₆	1.2
			C ₁ ...C ₇	7.6	F ₂ ...C ₅	1.0
			F ₂ ...C ₅	-7.6	C ₇ ...O ₈	0.9
					O ₆ ...C ₇	-0.9
3: 3-fluoropropanal 	C ₁	-6.9	F ₂ ...C ₈	-15.9	F ₂ ...H ₁₀	-1.7
			F ₂ ...O ₉	11.8	C ₁ ...F ₂	1.0
			C ₈ ...H ₁₀	7.4	C ₁ ...C ₅	-1.0
			C ₁ ...F ₂	6.5	C ₈ ...O ₉	0.9
			O ₉ ...H ₁₀	-5.7	F ₂ ...C ₈	-0.9
			C ₁ ...C ₈	5.6	C ₁ ...C ₈	0.8
					C ₈ ...H ₁₀	-0.7
4: N-(2-fluoroethyl)acetamide 	N ₆	-8.2	F ₂ ...N ₆	27.9	F ₂ ...N ₆	-2.5
	C ₁	-7.6	F ₂ ...C ₈	-16.5	H ₃ ...O ₉	2.4
			N ₆ ...C ₈	13.3	C ₁ ...C ₅	-2.2
			F ₂ ...H ₇	-12.8	H ₄ ...O ₉	-1.7
					C ₁ ...F ₂	1.3
					C ₅ ...N ₆	-1.2
5: 2-(fluoroethyl)isoindoline-1,3-dione 	C ₁	-3.9	F ₂ ...C ₂₁	-32.4	H ₃ ...O ₁₉	-2.6
	C ₂₁	3.7	F ₂ ...N ₂₀	32.2	F ₂ ...N ₂₀	-2.5
			F ₂ ...C ₁₈	-23.8	C ₁ ...C ₅	-1.7
			F ₂ ...O ₂₂	19.9	C ₁ ...N ₂₀	1.6
			C ₁ ...N ₂₀	-17.0	C ₁₈ ...O ₁₉	1.5

Table S2 (cont.)

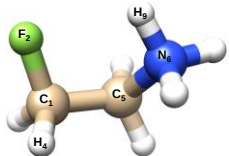
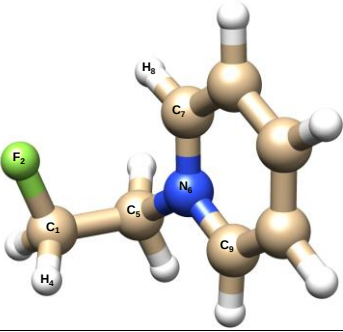
6: 2-fluoroethan-1-aminium 	C ₁ -28.2	F ₂ ...N ₆ 32.2 F ₂ ...H ₉ -26.6 C ₁ ...F ₂ 25.0 N ₆ ...H ₉ -15.2 C ₁ ...H ₉ 13.0	N ₆ ...H ₉ 6.6 C ₁ ...F ₂ 4.8 F ₂ ...N ₆ -4.4 C ₁ ...C ₅ -3.1 F ₂ ...H ₉ -2.4
7: 1-(2-fluoroethyl)pyridine-1-ium 	C ₁ -17.7	F ₂ ...N ₆ 34.3 F ₂ ...C ₇ 16.7 C ₁ ...F ₂ 14.6 C ₁ ...N ₆ -11.5	C ₁ ...F ₂ 3.0 C ₁ ...C ₅ -2.9 F ₂ ...C ₇ -2.3 F ₂ ...N ₆ -2.2

Table S3. Largest differences in the atomic energies (kcal/mol) between the F₂-C₁-C₅=O₆ *trans* and *cis* conformers in CH₂F-COX systems ($\Delta E = E_{trans} - E_{cis}$). Ball-and-stick models for the *trans* conformers including atom numbering.

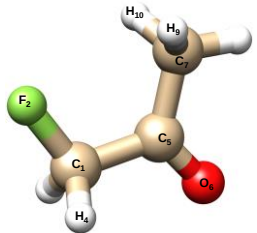
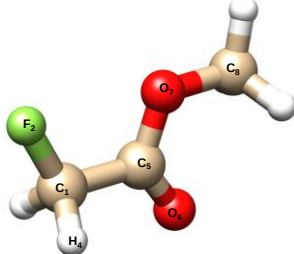
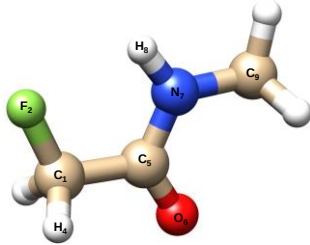
$\Delta E = E_{trans} - E_{cis}$	ΔE_{net}		$\Delta E_{int,clas}^{AB}$		$\Delta E_{int,xc}^{AB}$	
8: 1-fluoropropan-2-one 	C ₁	-17.2	C ₁ ...C ₅	-26.4	F ₂ ...O ₆	4.5
	C ₅	-10.3	C ₁ ...O ₆	24.9	F ₂ ...C ₇	-3.1
	F ₂	-5.2	F ₂ ...O ₆	-23.9	C ₁ ...O ₆	-3.0
			F ₂ ...C ₅	19.1	C ₁ ...F ₂	3.0
			C ₁ ...F ₂	17.6	C ₅ ...O ₆	3.0
					C ₁ ...C ₅	2.5
					C ₅ ...C ₇	-2.5
9: methyl-2-fluoroacetate 	C ₅	5.4	F ₂ ...O ₇	27.9	F ₂ ...O ₇	-5.2
	C ₁	-5.0	F ₂ ...O ₆	-25.2	C ₅ ...O ₆	4.6
	O ₇	4.9	C ₁ ...O ₆	18.2	F ₂ ...O ₆	4.2
			C ₁ ...O ₇	-14.7	C ₅ ...O ₇	-4.0
			C ₅ ...O ₇	-12.9	C ₁ ...O ₇	3.2
			F ₂ ...C ₈	-8.2	C ₁ ...O ₆	-2.9
10: 2-fluoro-N-methylacetamide 	C ₁	-23.5	F ₂ ...N ₇	39.6	N ₇ ...H ₈	8.3
	N ₇	15.5	C ₁ ...O ₆	28.6	C ₅ ...N ₇	-5.8
	H ₈	9.1	F ₂ ...O ₆	-25.1	F ₂ ...N ₇	-5.3
	F ₂	-6.9	N ₇ ...H ₈	-24.4	C ₅ ...O ₆	5.1
			F ₂ ...H ₈	-24.4	F ₂ ...O ₆	4.7
			C ₅ ...N ₇	-24.0	C ₁ ...F ₂	4.7
			C ₁ ...F ₂	23.4	C ₁ ...O ₆	-3.1
			C ₁ ...C ₅	-21.0	F ₂ ...H ₈	-3.0

Table S4. Largest differences in the atomic energies (kcal/mol) between the $g+/g+$ conformer of 1,2,3-trifluoropropane (**11** in Scheme 1) and the rest of conformers ($\Delta E = E_{g+/g+} - E_i$). Ball-and-stick models for the i -conformers including atom numbering.

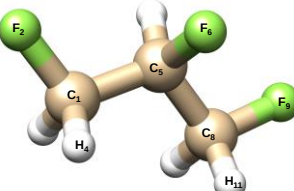
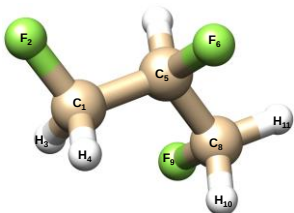
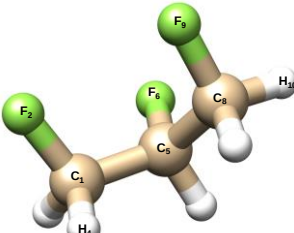
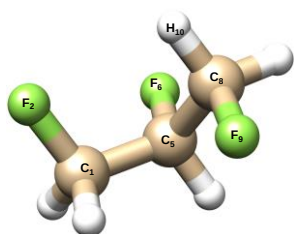
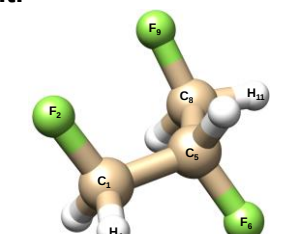
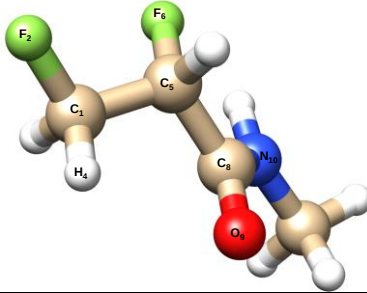
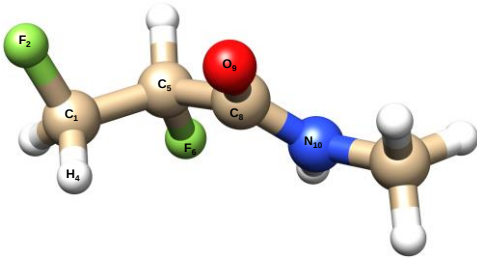
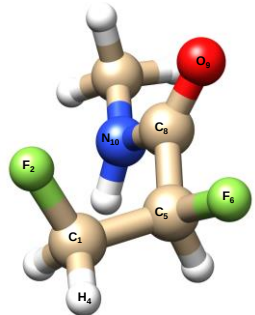
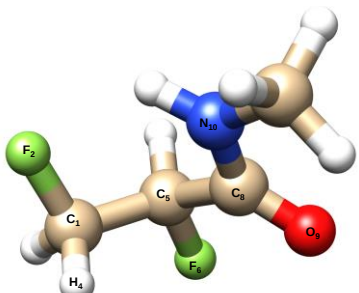
$\Delta E = E_{g+/g+} - E_i$	ΔE_{net}	$\Delta E_{int,clas}^{AB}$	$\Delta E_{int,xc}^{AB}$
$g+/g-$ 	C ₈ -6.4 C ₁ -2.6	C ₁ ...F ₉ -11.4 F ₂ ...F ₉ 6.7 C ₈ ...H ₁₀ 6.4 C ₈ ...F ₉ 6.3	C ₁ ...F ₉ -1.9 H ₄ ...F ₉ -1.7 C ₈ ...F ₉ 1.2 H ₃ ...H ₁₀ 1.0
$g+/anti$ 	H ₃ -1.4 H ₄ 1.4	F ₆ ...F ₉ 11.6 C ₅ ...F ₉ -7.5 F ₆ ...C ₈ -7.4	C ₅ ...C ₈ -2.8 H ₃ ...F ₉ 1.9 H ₄ ...F ₉ -1.6 C ₁ ...C ₅ 1.3
$g-/g+$ 	C ₈ -12.1 C ₁ -8.1 F ₂ -5.1 F ₉ -4.7	F ₂ ...C ₈ 19.2 F ₂ ...F ₉ -19.1 C ₁ ...C ₈ -14.2 C ₈ ...F ₉ 12.2 C ₁ ...F ₉ 10.5 C ₁ ...H ₄ 9.3	F ₂ ...F ₉ 3.6 C ₈ ...F ₉ 2.1 C ₁ ...F ₉ -1.9 C ₁ ...F ₂ 1.8 H ₄ ...F ₉ -1.7
$g-/anti$  F₂-C₁-C₅-F₆ -65.4 (-66.1) F₅-C₆-C₈-F₉ -174.9 (-174.9)	C ₁ 4.1	F ₂ ...C ₈ 13.0 F ₆ ...F ₉ 11.6 F ₂ ...F ₉ -11.1 C ₅ ...F ₉ -7.6 F ₆ ...C ₈ -6.8	C ₅ ...C ₈ -3.2 F ₂ ...H ₁₀ 2.3 F ₂ ...C ₈ 1.5 C ₁ ...C ₅ 1.3
$anti/anti$ 	C ₈ -8.1 C ₅ 4.4 C ₁ -3.8	F ₂ ...F ₉ -17.5 F ₂ ...C ₈ 16.7 F ₂ ...F ₆ 12.8 F ₆ ...F ₉ 11.9 C ₁ ...C ₈ -10.1 F ₂ ...C ₅ -9.7	C ₅ ...C ₈ -2.76 F ₂ ...F ₉ 2.56 C ₁ ...F ₉ -2.02 H ₄ ...F ₉ -1.69 C ₁ ...C ₅ -1.67

Table S5. Largest differences in the atomic energies (kcal/mol) between the **g+/trans** conformer of 2,3-difluoro-*N*-methylpropanamide (**12** in Scheme 1) and the rest of conformers ($\Delta E = E_{g+/trans} - E_i$). Ball-and-stick models for the *i*-conformers including atom numbering.

$\Delta E = E_{g+/trans} - E_i$	ΔE_{net}		$\Delta E_{int,clas}^{A \cdots B}$		$\Delta E_{int,xc}^{A \cdots B}$	
g-/trans 	C8	9.4	F2...C8	-38.4	H4...O9	3.1
			F2...N10	29.8	H3...O9	-2.0
			C1...C8	22.9	F2...C8	-1.0
			F2...O9	21.4	C5...C8	1.0
			C1...N10	-15.8		
anti/trans 	C1	-10.6	F2...N10	25.9	C1...C5	-4.0
	C8	5.4	F2...C8	-16.2	H3...O9	-2.1
			F2...F6	13.4	H3...F6	1.8
					C1...F2	1.8
					C8...N10	-1.6
g+/cis 	N10	19.9	F6...N10	40.2	N10...H11	9.2
	C5	-19.4	C8...N10	-32.9	C8...N10	-7.5
	H11	10.0	N10...H12	-28.2	C8...O9	7.0
	C1	-8.0	C5...O9	28.0	F6...N10	-5.5
					F6...O9	5.2
					C5...F6	4.3
anti/cis 	C5	-17.3	F6...N10	37.6	F6...O9	5.7
	C1	7.5	C5...O9	28.9	F6...N10	-5.5
			C1...O9	-28.6	C8...O9	5.0
			F6...O9	-27.1	C8...N10	-4.5
					C5...F6	4.1
					F2...H11	3.7
					C1...C5	-3.4

IQF analysis of fragments from an α,β -difluoro- γ -amino-acid

Figure S1. Ball-and-stick representation of the nine conformers optimized for 2-(2,3-difluoropropyl)isoindoline-1,3-dione (**13** in Scheme 1). F-C-C-F and F-C-C-N dihedral angles ($^\circ$) and selected interatomic distances ($\text{\AA}$) measured in the HF-D3/cc-pVTZ optimized structures (RI-MP2/cc-pVTZ in parentheses) are included. Energy differences ($E_{g^+/g^-} - E_i$) in kcal/mol computed at the HF-D3/cc-pVTZ, RI-MP2/cc-pVTZ (in parentheses) and DLPNO-CCSD(T)/aug-cc-pVTZ [in brackets] levels of theory are also included.

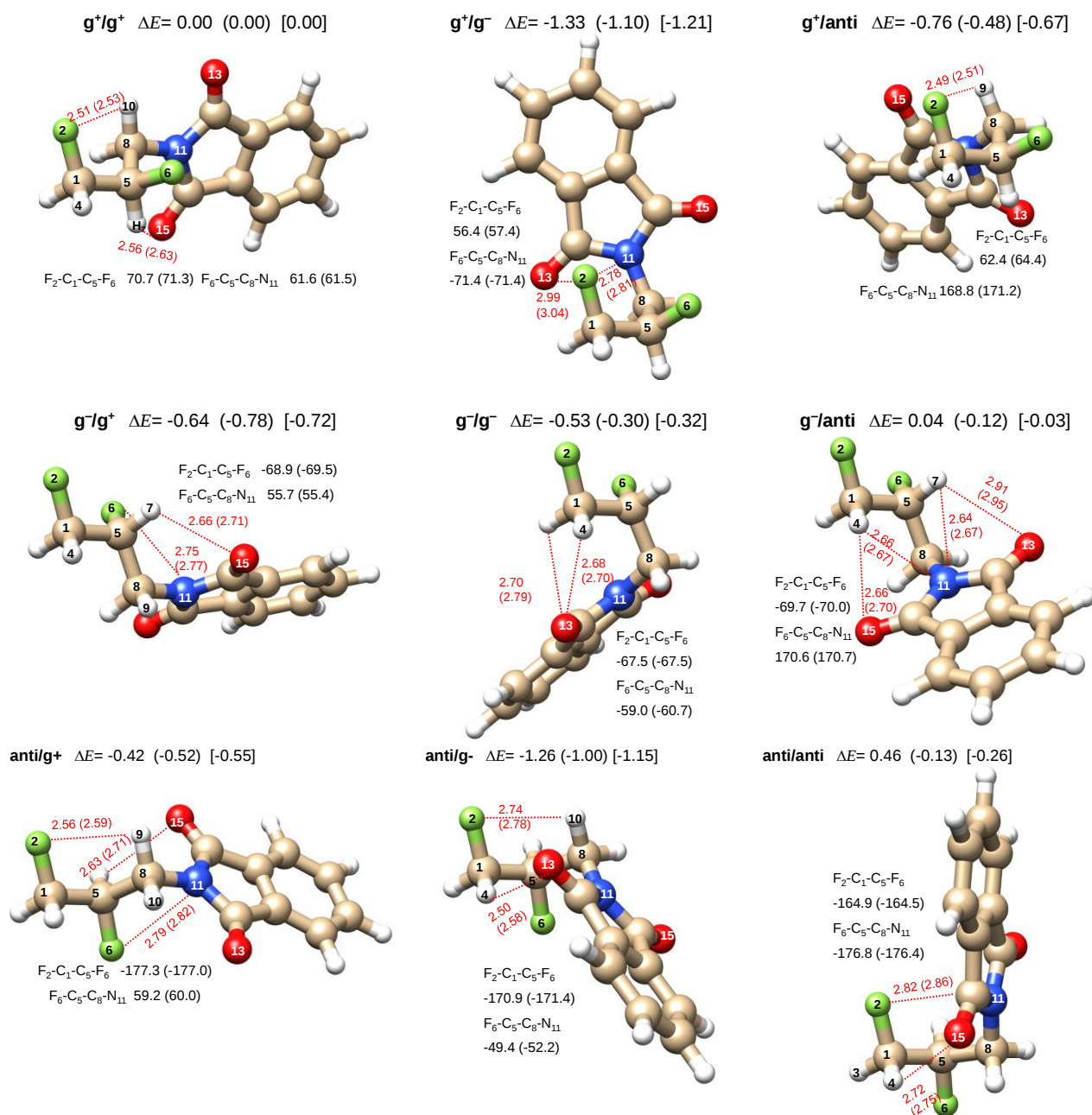
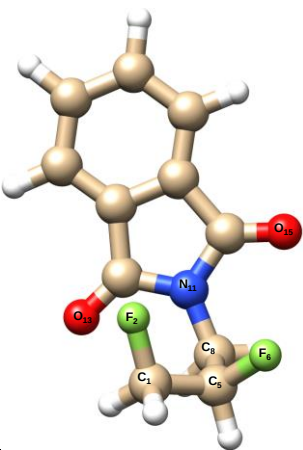
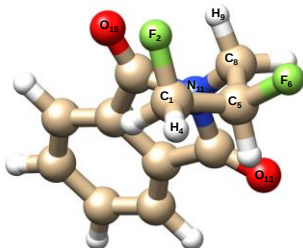
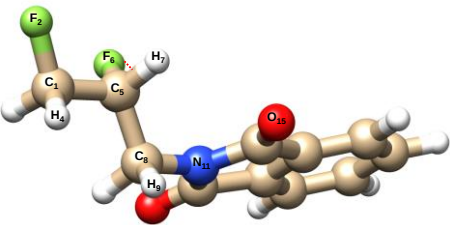
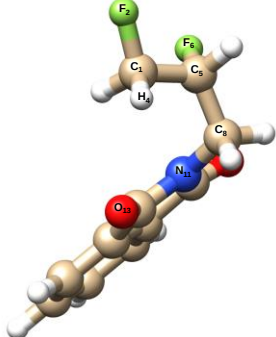


Table S6. IQF energy components at the HF-D3/cc-pVTZ level (in kcal/mol) for the energy differences among the nine conformers optimized for 2-(2,3-difluoropropyl)isoindoline-1,3-dione (**13**). A three-fragment partitioning scheme ($P=\text{CH}_2\text{F}$, $Q=\text{CHF}$, and $R=\text{CH}_2\text{isoindoline-1,3-dione}$) is assumed for the different conformers. Relative energies are given as $E_{g+/g+} - E_i$, so that a negative value means further stabilization of $g+/g+$.

	ΔE_{net}^P	ΔE_{net}^Q	ΔE_{net}^R	$\Delta E_{\text{int,disp}}$			$\Delta E_{\text{int,xc}}$			$\Delta E_{\text{int,class}}$			ΔE_{IQA}	$\Delta E_{\text{HF-}}$
				$P\cdots Q$	$P\cdots R$	$Q\cdots R$	$P\cdots Q$	$P\cdots R$	$Q\cdots R$	$P\cdots Q$	$P\cdots R$	$Q\cdots R$		
$\Delta E = E_{g+/g+} - E_{g+/g-}$	-7.2	3.1	-4.5	0.1,	2.3,	-1.1	-0.7,	8.6,	-4.4	0.1,	1.6,	-0.2	-2.2	-1.3
$\Delta E = E_{g+/g+} - E_{g+/anti}$	-2.3	2.1	0.2	0.0,	1.3,	-0.6	-0.0,	3.4,	-3.9	-0.2,	-0.9,	0.0	-1.1	-0.8
$\Delta E = E_{g+/g+} - E_{g-/g+}$	0.8	-1.5	1.8	0.0,	-0.1,	-0.1	0.3,	-1.4,	-0.4	-0.8,	-0.4,	0.0	-1.7	-0.6
$\Delta E = E_{g+/g+} - E_{g-/g-}$	-5.9	2.4	1.0	0.1,	1.8,	-1.0	0.4,	5.1,	-3.5	-0.3,	0.2,	-0.3	-0.1	-0.5
$\Delta E = E_{g+/g+} - E_{g-/anti}$	-3.8	1.6	1.7	0.0,	1.6,	-0.9	0.5,	3.7,	-4.6	-0.5,	-0.0,	0.0	-0.7	0.0
$\Delta E = E_{g+/g+} - E_{anti/g+}$	0.5	0.1	0.7	0.0,	-0.0,	-0.1	-2.4,	-0.4,	0.1	1.6,	-0.4,	-0.5	-0.9	-0.4
$\Delta E = E_{g+/g+} - E_{anti/g-}$	-4.4	2.1	-1.4	0.1,	1.6,	-0.9	-1.9,	5.2,	-3.1	2.0,	-0.8,	-0.4	-1.9	-1.3
$\Delta E = E_{g+/g+} - E_{anti/anti}$	-4.7	1.7	-2.7	0.0,	2.3,	-0.8	-2.8,	7.0,	-4.0	2.2,	1.7,	-0.3	-0.3	0.5

The optimized conformers for **13** alternate *gauche* ($g+$ and $g-$) and *anti* orientations in the adjacent $\text{F}_2\text{-C}_1\text{-C}_5\text{-F}_6$ and $\text{F}_6\text{-C}_5\text{-C}_8\text{-N}_{11}$ groups and remain quite close in energy (< 1.2 kcal/mol). The *gauche* effect between vicinal F atoms is reflected in favorable exchange-correlation and unfavorable electrostatic inter-fragment interactions ($\Delta E_{\text{int,xc}}^{P\cdots Q} = -2.4$ kcal/mol and $\Delta E_{\text{int,class}}^{P\cdots Q} = 1.6$ kcal/mol for the *anti/g+* conformer; see also *anti/g-* and *anti/anti* in Table S6). Most likely, this determines the relative stability of the *anti/g+* conformer with respect to $g+/g+$ ($\Delta E^{\text{IQA}} = -0.9$ kcal/mol). Actually, the similar stability of all the conformers indicates that the placement of the bulky $-\text{CH}_2\text{-isoindole-1,3-dione}$ group (fragment R) induces little or none conformational effects. This is not entirely unexpected because the presence of the vicinal methylene group and the relatively large size and *symmetric* structure of the isoindole moiety can yield similar $P\cdots R$ and $Q\cdots R$ interactions with the smaller $P=\text{CH}_2\text{F}$ and $Q=\text{CHF}$ fragments across the various conformers. This observation is not incompatible with the fact that the net and inter-fragment IQF energies (and the underlying atomic contributions) involving fragment R vary significantly upon conformational change. For example, on going from the extended $g+/g+$ reference to the more compact $g+/g-$ and *anti/anti* conformers, the closer $Q\cdots R$ contacts in $g+/g+$ are replaced by the $P\cdots R$ ones. These structural changes leave a mark on the $\Delta E_{\text{int,xc}}^{P\cdots R}$ values of 8.6 and 7.0 kcal/mol for $g+/g-$ and *anti/anti*, respectively, and on the $\Delta E_{\text{int,xc}}^{Q\cdots R}$ ones of -4.4 and -4.0 kcal/mol. Parallel energy variations of 2.3 and -1.0 kcal/mol arise in the dispersion $\Delta E_{\text{int,disp}}^{P\cdots R}$ and $\Delta E_{\text{int,disp}}^{Q\cdots R}$ terms. The $P\cdots R$ and $Q\cdots R$ interactions are also concomitant with opposite changes in the net fragment energies ($\Delta E_{\text{net}}^P + \Delta E_{\text{net}}^Q + \Delta E_{\text{net}}^R = -8.6$ and -5.7 kcal/mol for $g+/g-$ and *anti/anti*).

Table S7. Largest differences in the atomic energies (kcal/mol) between the **g+/g+** conformer of 2-(2,3-difluoropropyl)isoindoline-1,3-dione (**13** in Scheme 1) and the rest of conformers ($\Delta E = E_{g+/g+} - E_i$). Ball-and-stick models for the *i*-conformers including atom numbering.

$\Delta E = E_{g+/g+} - E_i$	ΔE_{net}		$\Delta E_{int,clas}^{AB}$		$\Delta E_{int,xc}^{AB}$	
g+/g- 	C ₁₄	-6.8	F ₂ ...C ₁₂	69.9	F ₂ ...N ₁₁	4.7
	C ₅	-5.4	C ₁ ...C ₁₂	-57.8	C ₁₂ ...O ₁₃	-3.8
	F ₂	-4.2	F ₂ ...N ₁₁	-51.3		
			F ₂ ...O ₁₃	-48.9		
			C ₁ ...O ₁₃	44.3		
g+/anti 	C ₁₂	7.0	F ₂ ...C ₁₄	36.2	C ₁₄ ...O ₁₅	3.4
	C ₁	-3.8	C ₁ ...C ₁₄	-35.0	H ₇ ...O ₁₅	-3.1
	C ₅	-3.6	F ₆ ...N ₁₁	32.8		
	C ₈	3.3	F ₂ ...O ₁₅	-32.6		
g-/g+ 	C ₁	-4.7	F ₂ ...C ₁₂	-13.8	F ₂ ...H ₁₀	-2.0
			F ₂ ...N ₁₁	11.9	F ₂ ...C ₈	-1.9
			F ₂ ...O ₁₃	11.5		
g-/g- 	C ₁₂	7.0	C ₁ ...C ₁₂	-29.8	C ₁₄ ...O ₁₅	3.9
	H ₁₀	3.2	C ₁ ...O ₁₃	28.2	N ₁₁ ...C ₁₄	-3.6
	N ₁₁	3.2	F ₆ ...C ₁₂	-16.4	C ₁₂ ...O ₁₃	-3.3
					H ₇ ...O ₁₅	-3.1

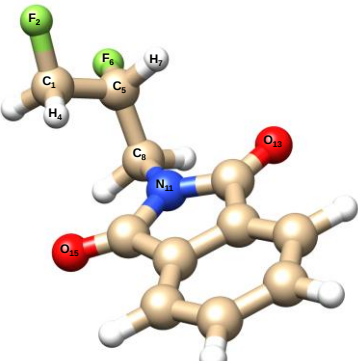
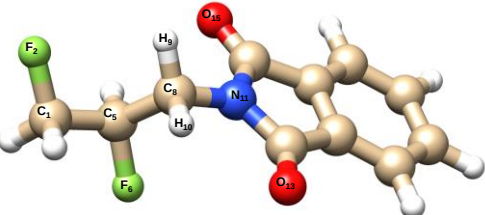
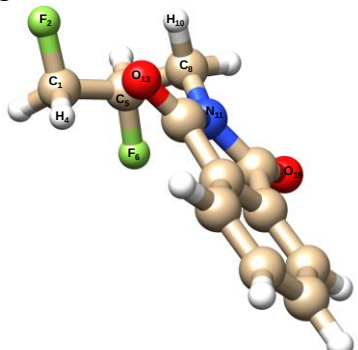
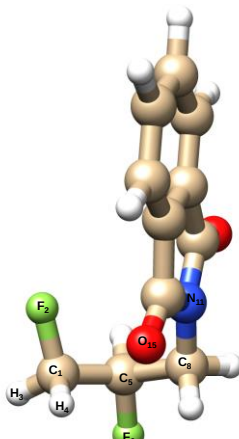
g-/anti 	C ₁₂ 6.0 H ₄ -3.0	F ₆ ⋯N ₁₁ 32.8 F ₆ ⋯C ₁₂ -29.5 F ₆ ⋯C ₁₄ -28.2 C ₁ ⋯C ₁₄ -23.8	H ₇ ⋯O ₁₅ -3.1 F ₆ ⋯N ₁₁ -2.6 H ₄ ⋯O ₁₅ 2.5
anti/g+ 	C ₁₄ -2.0 H ₁₀ 1.7	F ₂ ⋯F ₆ 11.9 F ₂ ⋯O ₁₅ -9.4 F ₂ ⋯O ₁₃ 8.2 F ₂ ⋯C ₁₄ 8.0 F ₂ ⋯C ₁₂ -7.9	C ₁ ⋯C ₅ -2.8 F ₂ ⋯H ₁₀ -2.0 F ₂ ⋯H ₉ 1.8
anti/g- 	N ₁₁ 6.4 C ₁₂ 4.7 C ₅ -4.0	C ₁ ⋯C ₁₂ -37.7 C ₁ ⋯O ₁₃ 36.1 F ₂ ⋯O ₁₃ -31.3 F ₂ ⋯C ₁₂ 30.3	H ₄ ⋯O ₁₃ 3.6 C ₁₄ ⋯O ₁₅ 3.4 C ₁ ⋯C ₅ -3.2 H ₇ ⋯O ₁₅ -3.1
anti/anti 	C ₁ 6.6 H ₁₀ 3.4 C ₁₄ -3.2	F ₂ ⋯C ₁₄ 66.7 F ₂ ⋯N ₁₁ -52.0 C ₁ ⋯C ₁₄ -46.9 F ₂ ⋯O ₁₅ -44.8	F ₂ ⋯N ₁₁ 4.5 C ₁ ⋯C ₅ -3.5

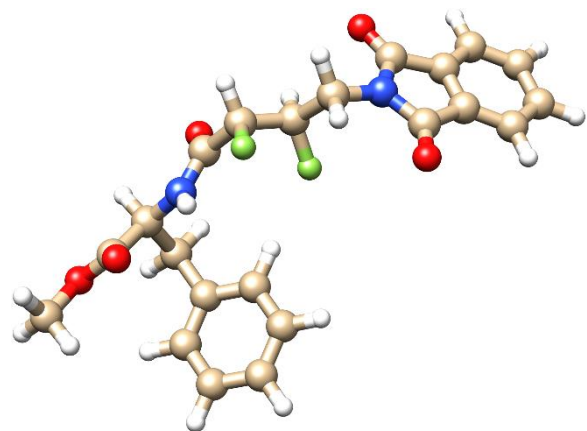
Table S8. Absolute (a.u) and relative $E_{14a-conf1}-E_{14a-confi}$ energies (kcal/mol) of the conformers optimize for **14a** at HF-D3/cc-pVTZ in gas-phase and in chloroform. Selected dihedral angles ($^{\circ}$) and intramolecular distances ($\text{\AA}$) are also included for the optimized structures (values in parentheses correspond to the solution structures). For the phenyl and the isoindoline-1,3-dione aromatic rings, the distance between the center of mass and the angle between the planes are also included.

	HF-D3/cc-pVTZ	HF-D3/cc-pVTZ SMD	F ₂₃ -C ₂₂ -C ₂₅ =O ₂₆	F ₂₃ -C ₂₂ -C ₁₉ -F ₂₀	F ₂₀ -C ₁₉ -C ₁₆ -N ₁₃	Aromatic rings
14a-conf1	-1527.900462 (0.00)	-1527.938920 (0.00)	-179.4 (-177.7)	67.5 (66.4)	61.1 (61.2)	8.8/48.0 (8.9/43.7)
14a-conf2	-1527.901636 (0.74)	-1527.937246 (-1.05)	171.2 (169.0)	-58.4 (-58.8)	53.6 (56.2)	7.4/35.6 (7.4/39.3)
14a-conf3	-1527.898920 (-0.97)	-1527.934012 (-3.08)	147.7 (147.1)	-78.1 (-76.8)	55.3 (52.0)	5.2/15.6 (5.3/17.3)
14a-conf4	-1527.898530 (-1.21)	-1527.935687 (-2.03)	-178.3 (-177.5)	57.6 (59.7)	166.9 (170.6)	10.3/79.8 (10.3/83.9)
14a-conf5	-1527.897323 (-1.97)	-1527.935517 (-2.14)	177.1 (171.9)	-143.2 (-159.7)	71.0 (67.3)	5.1/12.1 (5.2/4.2)
14a-conf6	-1527.896251 (-2.64)	-1527.932337 (-4.13)	1.8 (2.0)	-153.8 (-154.0)	-162.1 (-162.5)	8.2/87.0 (8.2/77.4)
14a-conf7	-1527.894583 (-3.69)	-1527.930825 (-5.08)	-178.2 (-177.3)	-55.6 (-49.3)	40.0 (50.0)	6.9/86.2 (6.7/71.8)
14a-conf8	-1527.894558 (-3.70)	-1527.931927 (-4.39)	-147.4 (-146.8)	-45.2 (-45.0)	-47.0 (-52.9)	9.0/61.4 (8.7/55.2)
14a-conf9	-1527.894194 (-3.93)	-1527.934486 (-2.78)	177.9 (-179.5)	57.7 (59.7)	167.2 (171.3)	10.3/55.3 (10.3/54.7)
14a-conf10	-1527.893735 (-4.22)	-1527.934880 (-2.54)	176.4 (179.4)	66.2 (65.0)	61.2 (61.6)	9.2/51.6 (9.4/58.8)
14a-conf11	-1527.893705 (-4.24)	-1527.932785 (-3.85)	169.4 (175.5)	49.6 (54.4)	167.4 (170.7)	8.1/75.1 (8.9/79.6)

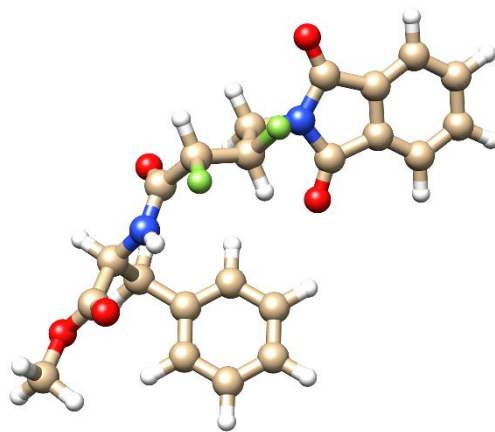
Table S8 (cont.)

	HF-D3/cc-pVTZ	HF-D3/cc-pVTZ SMD	F ₂₃ -C ₂₂ -C ₂₅ =O ₂₆	F ₂₃ -C ₂₂ -C ₁₉ -F ₂₀	F ₂₀ -C ₁₉ -C ₁₆ -N ₁₃	Aromatic rings
14a-conf12	-1527.893302 (-4.49)	-1527.932785 (-3.85)	169.4 (168.4)	-58.6 (-56.6)	56.3 (59.8)	8.5/85.5 (8.3/74.7)
14a-conf13	-1527.893188 (-4.56)	-1527.932512 (-4.02)	174.6 (176.5)	47.4 (49.8)	-66.0 (-64.9)	8.1/48.8 (8.7/46.1)
14a-conf14	-1527.892694 (-4.87)	-1527.934433 (-2.82)	172.8 (178.0)	54.9 (59.5)	167.7 (171.9)	8.6/50.9 (10.2/55.8)
14a-conf15	-1527.890338 (-6.35)	-1527.928668 (-6.43)	-10.8 (-11.7)	-165.7 (-167.4)	44.3 (44.4)	9.9/60.5 (10.0/61.0)
14a-conf16	-1527.889986 (-6.57)	-1527.929768 (-5.74)	-12.4 (-14.4)	58.0 (55.8)	-171.9 (-170.1)	7.2/88.0 (7.2/86.4)
14a-conf17	-1527.888970 (-7.21)	-1527.928708 (-6.41)	-21.5 (-22.0)	48.4 (48.5)	-177.2 (-174.5)	5.7/72.4 (5.8/66.5)
14a-conf18	-1527.888613 (-7.44)	-1527.929018 (-6.21)	-16.4 (-16.4)	53.2 (51.6)	-173.7 (-172.5)	6.9/73.3 (6.8/75.2)
14a-conf19	-1527.887910 (-7.88)	-1527.928883 (-6.30)	-17.1 (-16.6)	43.8 (44.0)	-174.2 (-173.5)	6.1/66.6 (6.1/69.4)
14a-conf20	-1527.887650 (-8.04)	-1527.929098 (-6.16)	-17.7 (-18.6)	47.5 (46.1)	-175.9 (-174.2)	6.2/64.9 (6.3/65.5)

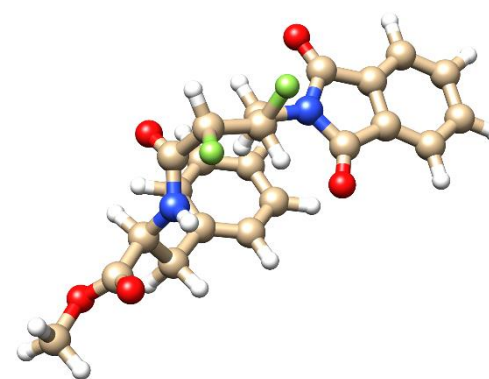
Figure S2. Ball-and-stick representation of the conformers optimized for **14a**.



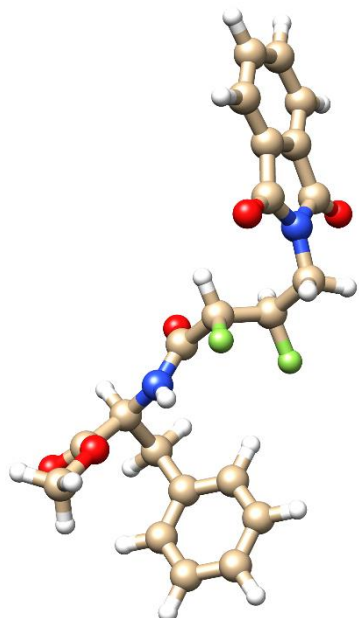
14a-conf1



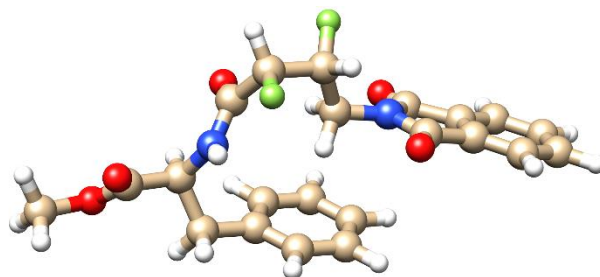
14a-conf2



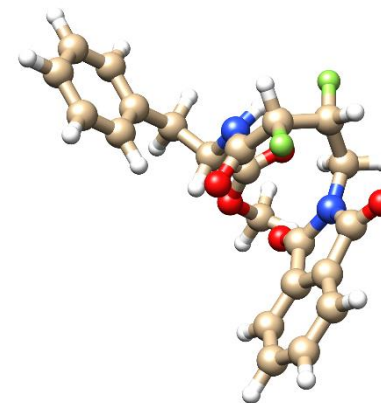
14a-conf3



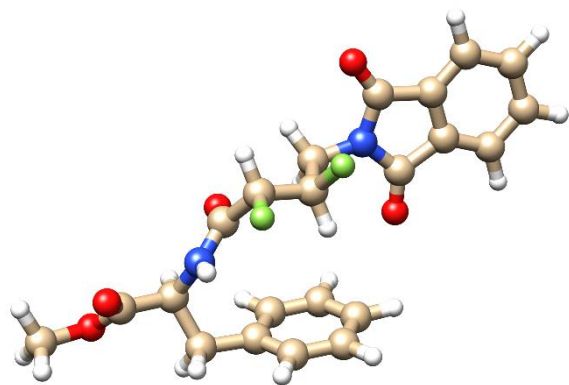
14a-conf4



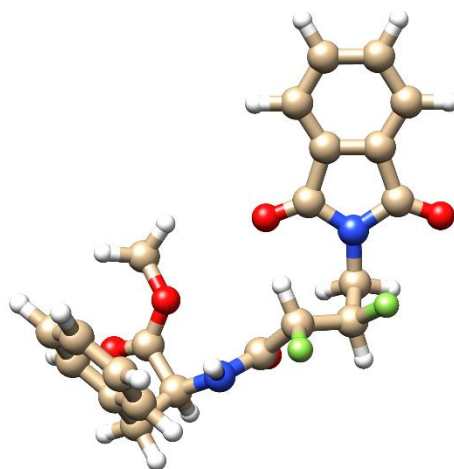
14a-conf5



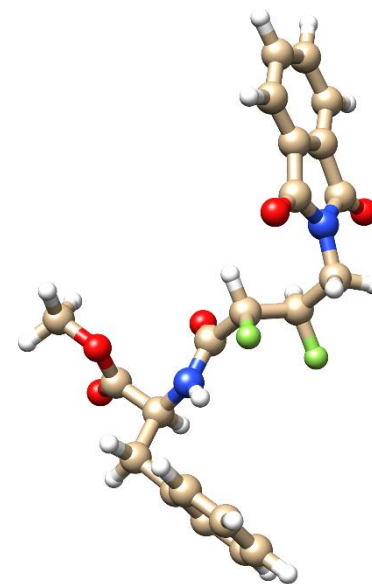
14a-conf6



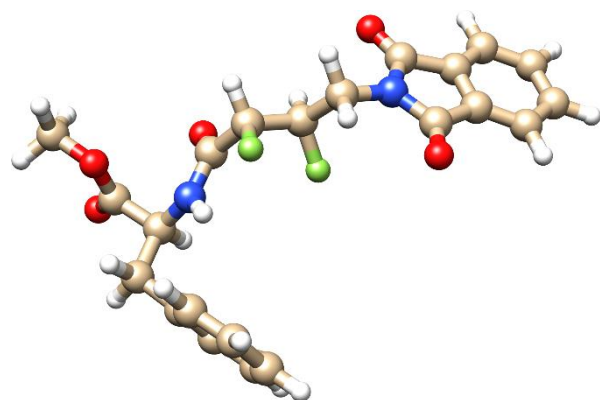
14a-conf7



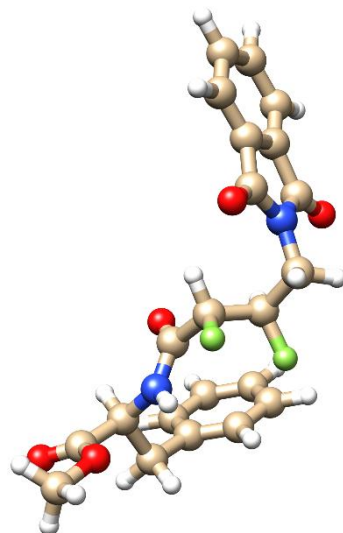
14a-conf8



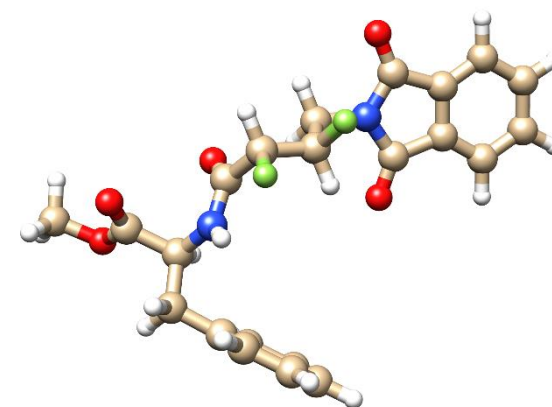
14a-conf9



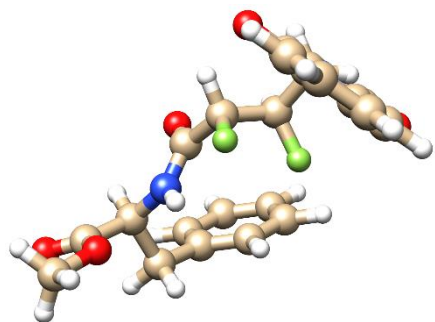
14a-conf10



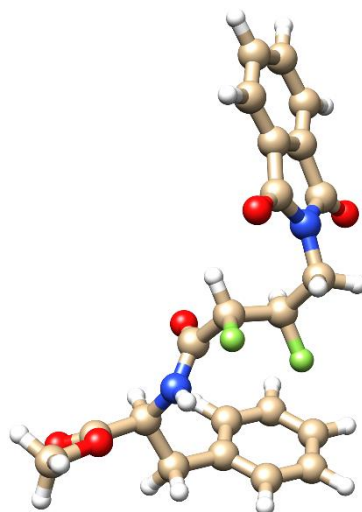
14a-conf11



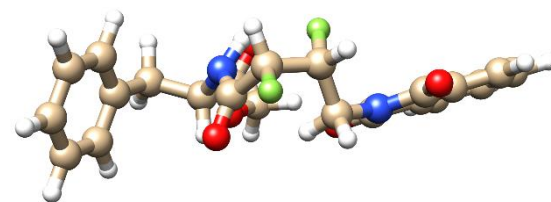
14a-conf12



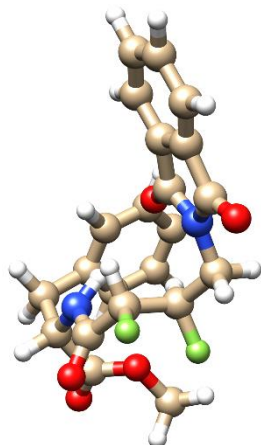
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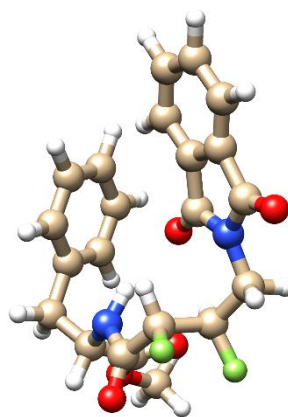
14a-conf14



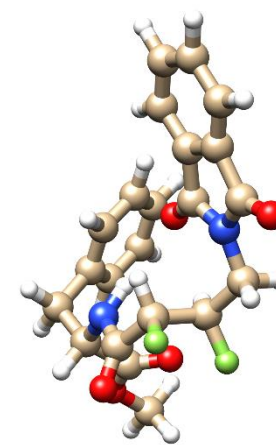
14a-conf15



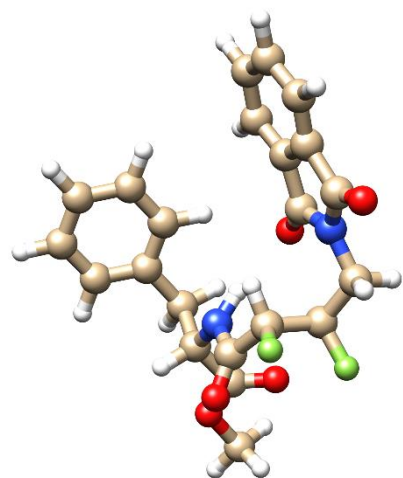
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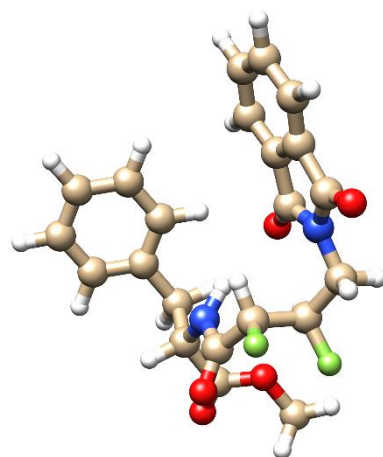
14a-conf17



14a-conf18



14a-conf19



14a-conf20

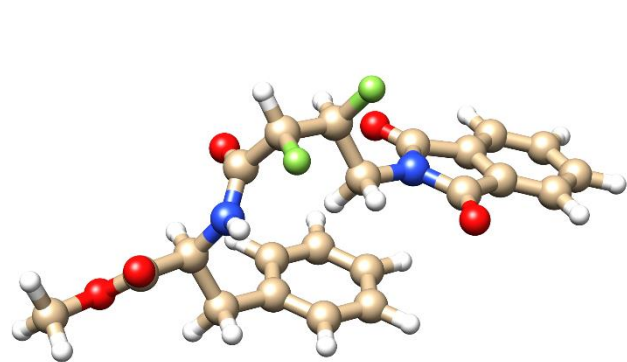
Table S9. Absolute (a.u) and relative $E_{14b-conf1}-E_{14b-confi}$ energies (kcal/mol) of the conformers optimize for **14b** at HF-D3/cc-pVTZ in gas-phase and in chloroform. Selected dihedral angles ($^{\circ}$) and intramolecular distances ($\text{\AA}$) are also included for the optimized structures (values in parentheses correspond to the solution structures). For the phenyl and the isoindoline-1,3-dione aromatic rings, the distance between the center of mass and the angle between the planes are also included.

	HF-D3/cc-pVTZ	HF-D3/cc-pVTZ SMD	F ₂₃ -C ₂₂ -C ₂₅ =O ₂₆	F ₂₃ -C ₂₂ -C ₁₉ -F ₂₀	F ₂₀ -C ₁₉ -C ₁₆ -N ₁₃	Aromatic rings
14b-conf1	-1527.902500 (0.00)	-1527.939296 (0.00)	178.6 (176.9)	-61.5 (-62.1)	-57.8 (-59.7)	5.1/9.4 (5.3/2.9)
14b-conf2	-1527.901093 (-0.88)	-1527.938076 (-0.76)	164.2 (163.8)	62.7 (62.8)	-59.5 (-57.6)	7.1/56.2 (7.2/55.6)
14b-conf3	-1527.900464 (-1.28)	-1527.937182 (-1.33)	-154.2 (-159.5)	-59.6 (-59.8)	65.2 (63.1)	6.0/36.2 (6.1/47.0)
14b-conf4	-1527.900032 (-1.55)	-1527.936010 (-2.06)	-161.5 (-164.3)	-60.5 (-60.9)	62.9 (62.0)	6.2/49.8 (6.2/52.7)
14b-conf5	-1527.896562 (-3.73)	-1527.933585 (-3.58)	151.5 (153.9)	-175.1 (178.2)	-63.9 (-62.4)	5.4/31.7 (6.2/29.4)
14b-conf6	-1527.895575 (-4.34)	-1527.931327 (-5.00)	173.3 (175.3)	62.7 (66.5)	166.5 (166.9)	9.7/60.8 (9.9/62.1)
14b-conf7	-1527.895159 (-4.61)	-1527.933710 (-3.50)	155.2 (152.6)	174.9 (171.8)	47.1 (54.3)	8.8/44.5 (9.3/54.2)
14b-conf8	-1527.894857 (-4.80)	-1527.932939 (-3.99)	-178.7 (173.3)	-63.3 (-65.2)	53.0 (51.4)	9.0/31.2 (9.1/31.2)
14b-conf9	-1527.894542 (-4.99)	-1527.935381 (-2.46)	-33.6 (-31.1)	71.8 (71.1)	-165.8 (-167.2)	7.5/47.1 (7.4/46.4)
14b-conf10	-1527.893634 (-5.56)	-1527.930076 (-5.78)	154.0 (153.0)	165.9 (166.6)	-178.9 (-176.8)	7.8/86.5 (7.7/86.6)
14b-conf11	-1527.892823 (-6.07)	-1527.933524 (-3.62)	176.2 (168.2)	-64.0 (-66.1)	52.1 (50.8)	9.2/88.7 (9.3/81.0)

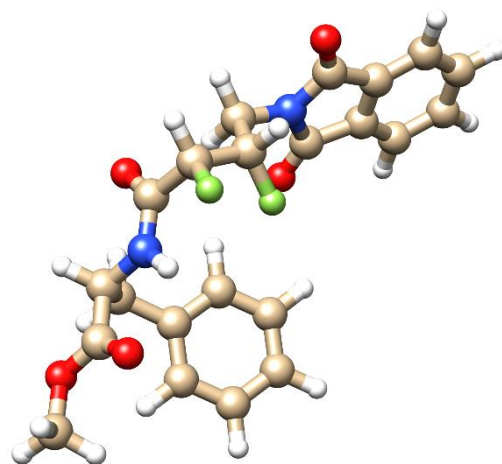
Table S9 (cont.)

	HF-D3/cc-pVTZ	HF-D3/cc-pVTZ SMD	F ₂₃ -C ₂₂ -C ₂₅ =O ₂₆	F ₂₃ -C ₂₂ -C ₁₉ -F ₂₀	F ₂₀ -C ₁₉ -C ₁₆ -N ₁₃	Aromatic rings
14b-conf12	-1527.892659 (-6.18)	-1527.932022 (-4.56)	-163.2 (160.9)	-149.5 (-176.1)	-52.0 (-66.5)	7.0/77.7 (5.3/39.6)
14b-conf13	-1527.892565 (-6.23)	-1527.930652 (-5.42)	158.8 (160.3)	-172.8 (-177.0)	-67.9 (-66.9)	4.9/41.4 (5.3/39.5)
14b-conf14	-1527.891884 (-6.66)	-1527.929878 (-5.91)	23.8 (21.6)	-63.2 (-61.4)	-78.1 (-77.9)	5.6/77.5 (5.8/74.6)
14b-conf15	-1527.890527 (-7.51)	-1527.929359 (-6.24)	-164.4 (-168.4)	-150.5 (-157.5)	-51.5 (-54.9)	6.9/78.9 (6.9/87.6)
14b-conf16	-1527.890134 (-7.76)	-1527.931522 (-4.88)	-23.7 (-15.4)	-71.5 (-69.7)	-67.2 (-83.9)	10.5/62.5 (9.1/84.0)
14b-conf17	-1527.888887 (-8.54)	-1527.928227 (-6.94)	165.5 (163.2)	-177.5 (179.9)	44.2 (51.1)	7.8/66.4 (8.0/77.2)
14b-conf18	-1527.888717 (-8.65)	-1527.928716 (-6.64)	-68.1 (-41.1)	176.8 (172.6)	45.9 (53.1)	8.8/86.2 (7.8/61.8)
14b-conf19	-1527.886233 (-10.21)	-1527.926181 (-8.23)	151.4 (152.5)	161.5 (164.2)	179.1 (-179.4)	11.2/57.8 (11.2/61.8)
14b-conf20	-1527.885198 (-10.86)	-1527.926237 (-8.19)	178.1 (161.1)	-148.2 (-176.8)	-56.1 (-59.3)	12.0/50.1 (11.9/86.0)

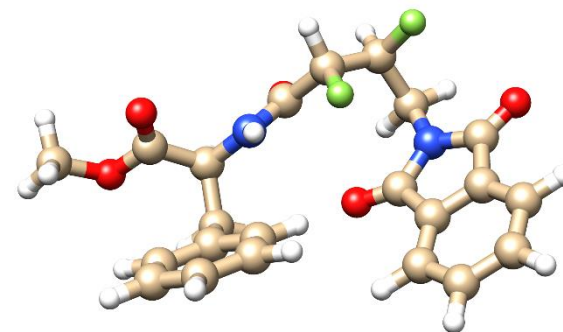
Figure S3. Ball-and-stick representation of the conformers optimized for **14b**.



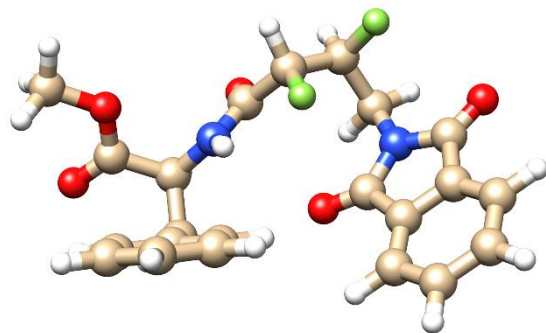
14b-conf1



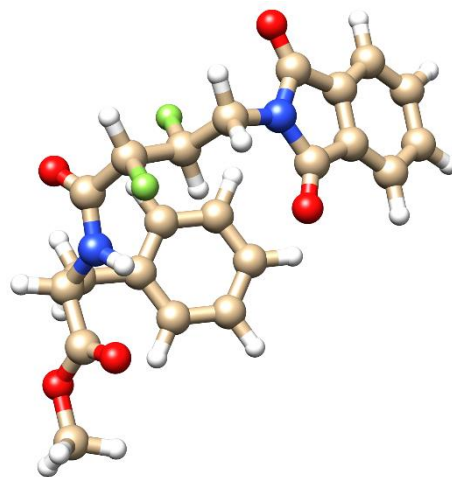
14b-conf2



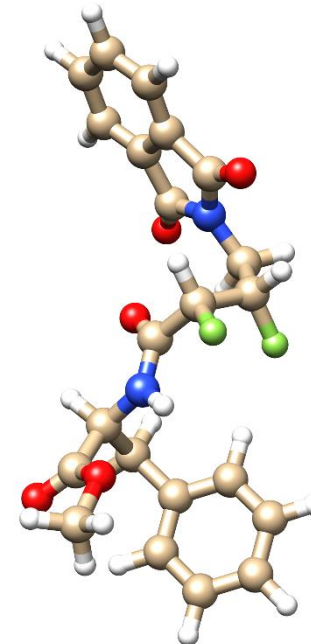
14b-conf3



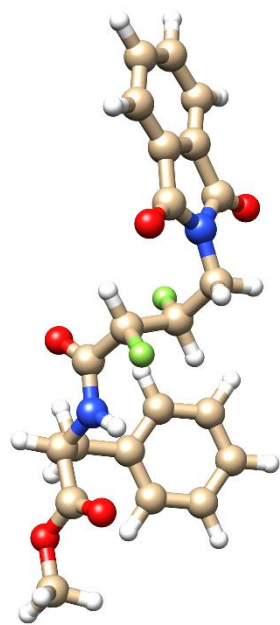
14b-conf4



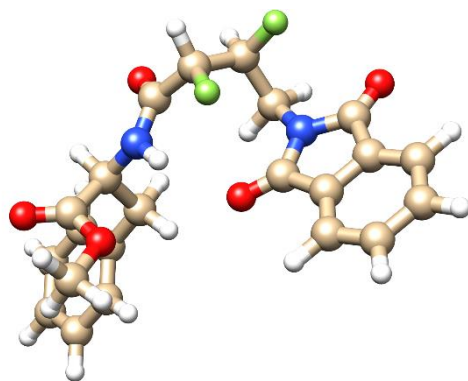
14b-conf5



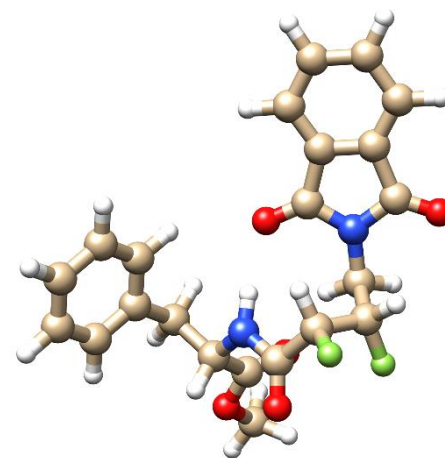
14b-conf6



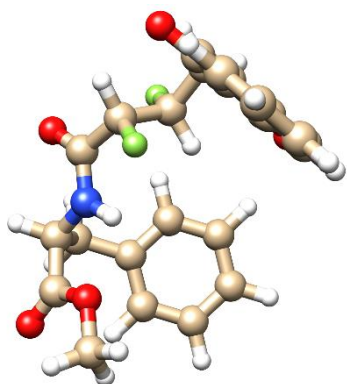
14b-conf7



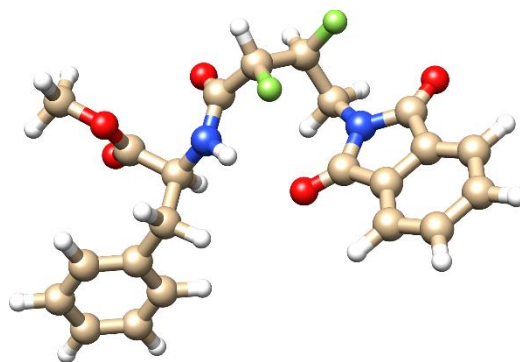
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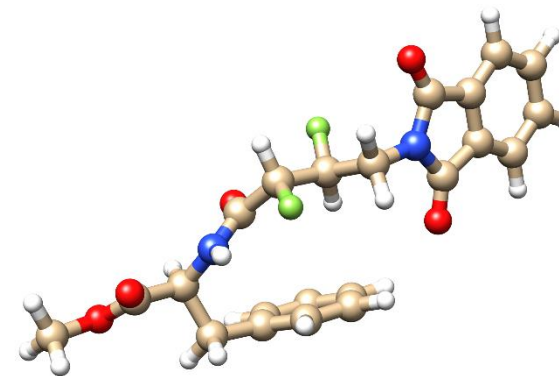
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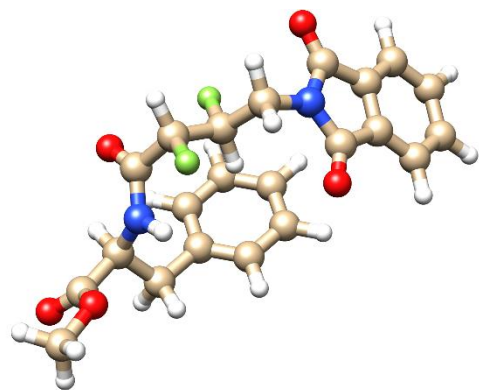
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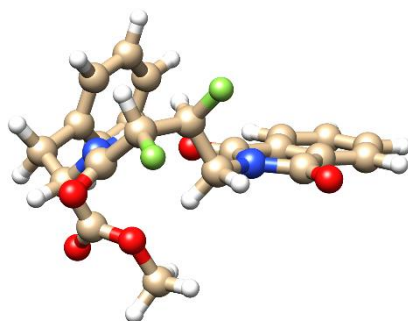
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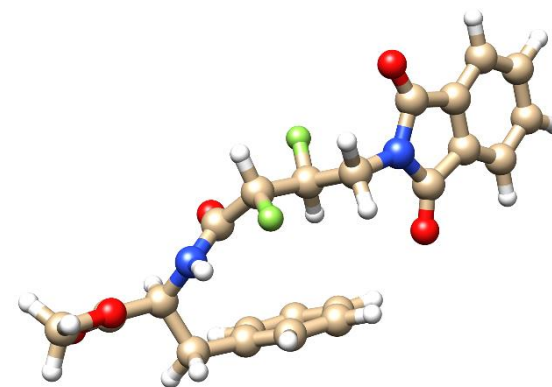
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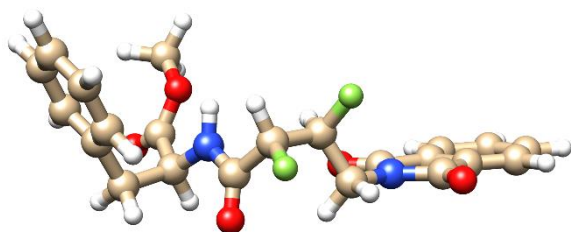
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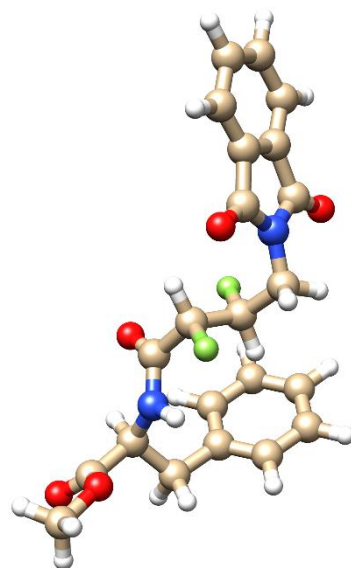
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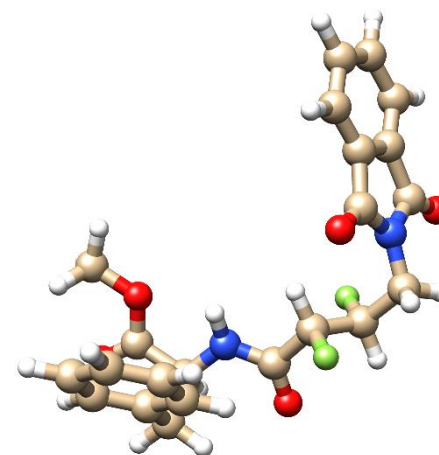
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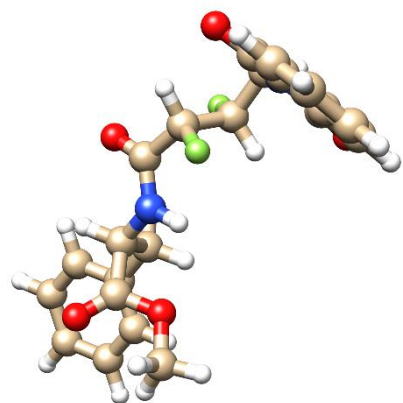
14b-conf16



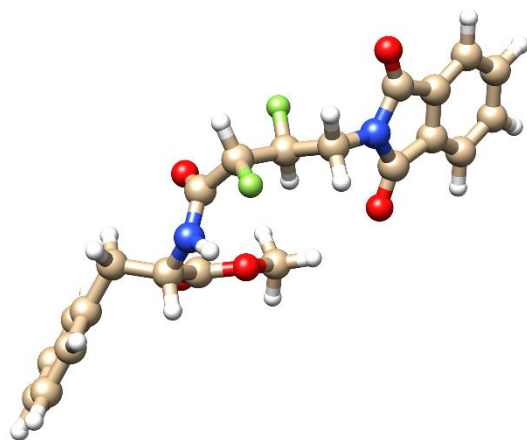
14b-conf17



14b-conf18



14b-conf19



14b-conf20

Table S10. Estimated NMR coupling constants (*Hz*) between vicinal fluorine and hydrogen atoms around the C₂₂-C₁₉ bond in selected conformers of the **14a** and **14b** compounds. Relative population of the different conformers at 300 K according to the Boltzmann distribution and the HF-D3/cc-pVTZ relative energies in chloroform. Coupling constants (*Hz*) obtained from the NMR spectra of **14a** and **14b** recorded in CDCl₃ at 300 K.

	$^3J_{FH}$	$^3J_{HF}$	P_i		$^3J_{FH}$	$^3J_{HF}$	P_i
14a-conf1	30.8	31.8	0.807	14b-conf1	30.8	8.1	0.704
14a-conf2	8.4	8.8	0.139	14b-conf2	6.6	31.7	0.197
14a-conf3	17.2	14.6	0.027	14b-conf3	31.3	10.4	0.076
14a-conf4	30.6	31.8	0.022	14b-conf4	31.6	10.5	0.022
14a-conf5	21.2	17.1	0.005	14b-conf5	5.2	3.9	0.002
Averaged	27.4	28.2	1.0	Averaged	26.1	13.0	1.0
NMR	27.0	28.9	--	NMR	23.0	21.4	--

Table S11. Largest differences in the atomic energies (kcal/mol) between the **14a-conf1** conformer and selected *i*-conformers.

$\Delta E = E_{14a-conf1} - E_i$	ΔE_{net}		$\Delta E_{int,clas}^{A...B}$		$\Delta E_{int,xc}^{A...B}$	
14a-conf2	C ₂₅	11.4	F ₂₀ ...C ₂₅	-43.9	H ₁₇ ...O ₂₆	5.0
	H ₁₇	-4.6	F ₂₀ ...O ₂₆	29.2	F ₂₀ ...H ₄₄	-3.2
	N ₂₇	-4.6	N ₁₃ ...O ₂₆	-28.3	C ₁₆ ...O ₂₆	3.1
	C ₁₉	-4.2	F ₂₀ ...N ₂₇	28.0	O ₁₂ ...H ₄₂	3.0
	C ₁₆	3.8	C ₁₉ ...C ₂₅	27.1	C ₁₁ ...O ₁₂	-2.3
	O ₁₂	-3.7	O ₁₂ ...N ₂₇	-25.6	O ₁₂ ...H ₄₄	2.3
	O ₂₆	-3.5	C ₁₁ ...N ₂₇	23.8	H ₂₁ ...O ₂₆	-2.3
	C ₂₂	-2.1	O ₁₂ ...C ₄₅	23.0	H ₁₈ ...F ₂₃	-2.0
			N ₁₃ ...C ₂₅	22.7	C ₁₉ ...C ₂₂	-1.9
			C ₁₁ ...C ₂₅	-22.5	C ₁₆ ...F ₂₃	-1.8
			C ₁₄ ...O ₂₆	21.7	F ₂₀ ...F ₂₃	1.8
14a-conf5	C ₂₅	14.5	N ₁₃ ...C ₂₅	33.2	N ₂₇ ...H ₂₈	6.3
	N ₂₇	10.6	N ₁₃ ...N ₂₇	-29.6	C ₁₉ ...C ₂₂	-5.5
	C ₁₉	-10.1	C ₁₄ ...C ₂₅	-28.3	H ₁₈ ...O ₂₆	3.6
	C ₁₁	-6.6	C ₁₄ ...N ₂₇	28.3	N ₁₃ ...C ₁₄	3.4
	H ₂₈	6.4	F ₂₀ ...N ₂₇	27.0	O ₁₅ ...H ₁₈	-3.4
	H ₁₇	-5.9	N ₁₃ ...O ₂₆	-26.5	O ₁₂ ...H ₁₈	3.3
	O ₂₆	-5.8	C ₂₅ ...N ₂₇	-25.9	F ₂₀ ...H ₄₄	-3.3
			F ₂₀ ...C ₂₅	-24.4	C ₂₅ ...N ₂₇	-3.0
			O ₁₅ ...N ₂₇	-23.9	O ₂₆ ...H ₃₃	-2.6
			O ₁₅ ...C ₂₅	23.7	O ₂₆ ...H ₃₀	2.6
			C ₁₁ ...N ₂₇	20.0	N ₁₃ ...C ₃₉	2.6
14a-conf6	N ₂₇	20.0	C ₁₁ ...C ₄₅	-194.5	N ₂₇ ...H ₂₈	8.0
	C ₂₅	16.5	O ₁₂ ...C ₄₅	194.1	F ₂₃ ...O ₂₆	5.9
	O ₂₆	-13.2	C ₁₁ ...C ₂₅	-172.0	F ₂₃ ...N ₂₇	-5.7
	C ₂₂	-12.7	C ₁₁ ...O ₄₆	136.5	C ₁₉ ...C ₂₂	-4.5
	O ₁₂	-11.0	N ₁₃ ...C ₂₅	136.2	H ₁₇ ...O ₄₆	4.3
	H ₂₈	8.6	C ₁₁ ...O ₂₆	135.6	O ₂₆ ...H ₄₄	4.2
	H ₁₇	-8.5	O ₁₂ ...O ₄₆	-130.9	C ₄₅ ...O ₄₆	-4.0
	C ₄₅	-8.4	O ₁₂ ...O ₄₇	-129.1	C ₂₅ ...N ₂₇	-4.0
	F ₂₃	-7.3	C ₁₁ ...N ₂₇	127.5	O ₁₂ ...O ₂₆	3.9
			O ₁₂ ...C ₂₅	126.4	C ₂₉ ...C ₃₁	3.7
			C ₁₁ ...O ₄₇	125.9	C ₁₆ ...H ₁₇	-3.4

Table S12. Equivalence between the filenames with the Cartesian coordinates for the HF/MP2 minima available as Supporting Information and the names of the compounds used in the text/Tables/Figures.

Filename	Compound	Filename	Compound
1_gauche_HF.xyz 1_gauche_MP2.xyz	1: 1,2-difluoroethane in gauche	1_anti_HF.xyz 1_anti_MP2.xyz	1: 1,2-difluoroethane in anti
2_gauche_HF.xyz 2_gauche_MP2.xyz	2: 2-fluoroethyl-acetate in gauche	2_anti_HF.xyz 2_anti_MP2.xyz	2: 2-fluoroethyl-acetate in anti
3_gauche_HF.xyz 3_gauche_MP2.xyz	3: 3-fluoropropanal in gauche	3_anti_HF.xyz 3_anti_MP2.xyz	3: 3-fluoropropanal in anti
4_gauche_HF.xyz 4_gauche_MP2.xyz	4: <i>N</i> -(2-fluoroethyl)acetamide in gauche	4_anti_HF.xyz 4_anti_MP2.xyz	4: <i>N</i> -(2-fluoroethyl)acetamide in anti
5_gauche_HF.xyz 5_gauche_MP2.xyz	5: 2-(2-fluoroethyl)isoindole-1,3-dione in gauche	5_anti_HF.xyz 5_anti_MP2.xyz	5: 2-(2-fluoroethyl)isoindole-1,3-dione in anti
6_gauche_HF.xyz 6_gauche_MP2.xyz	6: 2-fluoroethan-1-ammonium in gauche	6_anti_HF.xyz 6_anti_MP2.xyz	6: 2-fluoroethan-1-ammonium in anti
7_gauche_HF.xyz 7_gauche_MP2.xyz	7: 1-(2-fluoroethyl)pyridin-1-ium in gauche	7_anti_HF.xyz 7_anti_MP2.xyz	7: 1-(2-fluoroethyl)pyridine-1-ium in anti
8_trans_HF.xyz 8_trans_MP2.xyz	8: 1-fluoropropan-2-one in trans	8_cis_HF.xyz 8_cis_MP2.xyz	8: 1-fluoropropan-2-one in cis
9_trans_HF.xyz 9_trans_MP2.xyz	9: methyl-2-fluoroacetate in trans	9_cis_HF.xyz 9_cis_MP2.xyz	9: methyl-2-fluoroacetate in cis
10_trans_HF.xyz 10_trans_MP2.xyz	10: 2-fluoro- <i>N</i> -methylacetamide in trans	10_cis_HF.xyz 10_cis_MP2.xyz	10: 2-fluoro- <i>N</i> -methylacetamide in cis
11_g1g1_HF.xyz 11_g1g1_MP2.xyz	11: 1,2,3-trifluoropropane in g+/g+	11_g1g2_HF.xyz 11_g1g2_MP2.xyz	11: 1,2,3-trifluoropropane in g+/g-
11_g1anti_HF.xyz 11_g1anti_MP2.xyz	11: 1,2,3-trifluoropropane in g+/anti	11_g2g1_HF.xyz 11_g2g1_MP2.xyz	11: 1,2,3-trifluoropropane in g-/g+
11_g2anti_HF.xyz 11_g2anti_MP2.xyz	11: 1,2,3-trifluoropropane in g-/anti	11_antianti_HF.xyz 11_antianti_MP2.xyz	11: 1,2,3-trifluoropropane in anti/anti
12_g1trans_HF.xyz 12_g1trans_MP2.xyz	12: 2,3-difluoro- <i>N</i> -methylpropanamida in g+/trans	12_g2trans_HF.xyz 12_g2trans_MP2.xyz	12: 2,3-difluoro- <i>N</i> -methylpropanamida in g-/trans
12_antitrans_HF.xyz 12_antitrans_MP2.xyz	12: 2,3-difluoro- <i>N</i> -methylpropanamida in anti/trans	12_g1cis_HF.xyz 12_g1cis_MP2.xyz	12: 2,3-difluoro- <i>N</i> -methylpropanamida in g+/cis
12_anticis_HF.xyz 12_anticis_MP2.xyz	12: 2,3-difluoro- <i>N</i> -methylpropanamida in anti/cis	13_g1g1_HF.xyz 13_g1g1_MP2.xyz	13: 2-(2,3-difluoropropyl)isoindoline-1,3-dione in g+/g+
13_g1g2_HF.xyz 13_g1g2_MP2.xyz	13: 2-(2,3-difluoropropyl)isoindoline-1,3-dione in g+/g-	13_g1anti_HF.xyz 13_g1anti_MP2.xyz	13: 2-(2,3-difluoropropyl)isoindoline-1,3-dione in g+/anti
13_g2g1_HF.xyz 13_g2g1_MP2.xyz	13: 2-(2,3-difluoropropyl)isoindoline-1,3-dione in g+/g+	13_g2g2_HF.xyz 13_g2g2_MP2.xyz	13: 2-(2,3-difluoropropyl)isoindoline-1,3-dione in g-/g-
13_g2anti_HF.xyz 13_g2anti_MP2.xyz	13: 2-(2,3-difluoropropyl)isoindoline-1,3-dione in g-/anti	13_antig1_HF.xyz 13_antig1_MP2.xyz	13: 2-(2,3-difluoropropyl)isoindoline-1,3-dione in anti/g+
13_antig2_HF.xyz 13_antig2_MP2.xyz	13: 2-(2,3-difluoropropyl)isoindoline-1,3-dione in anti/g-	13_antianti_HF.xyz 13_antianti_MP2.xyz	13: 2-(2,3-difluoropropyl)isoindoline-1,3-dione in anti/anti
14a_conf1_HF.xyz	14a-conf1	14a_conf2_HF.xyz	14a-conf2
14a_conf5_HF.xyz	14a-conf5	14a_conf6_HF.xyz	14a-conf6

Table S13. IQF energy components for the energy difference (kcal/mol) between the *gauche/anti* conformers analyzed for 1,2-difluoroethane at different levels of theory. Two-fragment partitioning scheme ($P=\text{CH}_2\text{F}-$ and $Q=-\text{CH}_2\text{F}$) is assumed for the system. Geometrical changes in distances (Å) and angles ($^\circ$) between both conformers (*gauche-anti*) are also included.

$\Delta E = E_{\text{gauche}} - E_{\text{anti}}$	Δq^P	ΔE_{net}^P	ΔE_{net}^Q	$\Delta E_{\text{int,disp}}$	$\Delta E_{\text{int,xc}}$	$\Delta E_{\text{int,class}}$	$\Delta E_{\text{IQA}} (\Delta E)^{(a)}$	C-C	C-F	F-C-C	IQA wall time ^(b)
HF-D3/cc-pVTZ	0.00	0.6	0.4	0.0	-3.5	2.1	-0.5 (-0.3)	-0.009	-0.002	2.2	24
B3LYP-D3/cc-pVTZ ^(c)	0.00	1.2	1.4	0.0	-5.8	2.2	-1.0 (-0.9)	-0.013	-0.002	2.5	26
MP2/cc-pVTZ ^(d)	0.00	0.3	0.3	-	-4.2	2.4	-1.2 (-0.8)	-0.012	-0.002	2.3	1348

- (a) ΔE_{IQA} =IQA-reconstructed energy difference; ΔE = Exact energy difference at the corresponding level of theory. The difference between ΔE_{IQA} and ΔE is due to the numerical errors in the IQA calculations.
- (b) Wall time in min of the IQA calculations running on a 12-core Xeon platform.
- (c) The total energy of the Khon-Sham DFT energy is recovered with IQA using the scaling technique as described in Francisco, E.; Casals-Sainz, J. L.; Rocha-Rinza, T. and Martín Pendás, A. *Theoretical Chemistry Accounts* **2016**, 135.
- (d) The MP2 correlation energy is expressed as $E_{\text{corr}} = \sum_{iajb} t_{ij}^{ab} (ia|jb)$ where t_{ij}^{ab} are the MP2 amplitudes, $(ia|jb)$ are two electron integrals in the molecular

orbital basis (MO) (Mulliken convention) and i, j and a, b refer to occupied and virtual orbitals, respectively. The t_{ij}^{ab} amplitudes are obtained using a locally modified copy of the PySCF code by performing a single-point MP2/cc-pVTZ calculation on the corresponding optimized geometry. The MP2-correlated IQA calculations are also performed with the PROMOLDEN code. For further details on the correlated-IQA calculations see Casals-Sainz, J.L. Costales-Castro, A; Francisco, E. and Martín-Pendás, A. *Molecules* **2019**, 24, 2204. For PYSCF see: Sun, Q.; Berkelbach, T.C.; Blunt, N.S.; Booth, G.H.; Guo, S.; Li, Z.; Liu, J.; McClain, J.D.; Sayfutyarova, E.R.; Sharma, S.; et al. *PySCF: The Python-based simulations of chemistry framework*. *Wiley Interdiscip. Rev. Comput. Mol. Sci.* **2018**, 8, e1340.

Short comment on the results of Table S13

The three levels of theory predict similar geometrical changes upon the *anti*→*gauche* conformational change. The classical electrostatic penalty of the *gauche* conformation ($\Delta E_{\text{int,class}} \sim 2.1\text{-}2.4$ kcal/mol) is also quite similar among the three computational approaches. They coincide again in predicting a significant quantum mechanical stabilization ($\Delta E_{\text{int,xc}}$) of the *gauche* conformer, although the B3LYP-based energy (-5.8 kcal/mol) is significantly larger in absolute value than the MP2/HF ones (-4.2/-3.5). B3LYP-D3 also differs in the relative importance of the fragment net energy changes ($\Delta E_{\text{net}}^P, \Delta E_{\text{net}}^Q > 1$ kcal/mol), the HF-D3 and MP2 values being closer to each other. Overall, we see that, with respect to HF-D3, inclusion of the MP2 dynamical correlation does not alter the qualitative changes of the IQF energy terms and has only a small quantitative impact. Note also that the MP2-IQA calculations are computationally very demanding ($\sim 10^2$ more expensive than HF-IQA) so that they cannot be performed on large molecular systems.