

Supporting Information:

On the Applicability of Functional-Group Symmetry-Adapted Perturbation Theory and Other Partitioning Models for a Chiral Recognition – the Case of Popular Drug Molecules Interacting with Chiral Phases

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List of Figures

FS1 Simulated B97-D3/aug-cc-pVTZ IR spectra for the monomers under study.	4
FS2 Simulated B97-D3/aug-cc-pVTZ VCD spectra for complexes with phenethylamine.	4
FS3 Simulated B97-D3/aug-cc-pVTZ VCD spectra for complexes with proline.	5
FS4 Simulated B97-D3/aug-cc-pVTZ VCD spectra for the monomers under study.	5

List of Tables

TS1 B97-D3/aug-cc-pVTZ optimized geometry of the (S) Bac – (S) Phe complex.	6
TS2 B97-D3/aug-cc-pVTZ optimized geometry of the (S) Bac – (S) Pro complex.	7

TS3 B97-D3/aug-cc-pVTZ optimized geometry of the (R) Ibu – (R) Phe complex.	8
TS4 B97-D3/aug-cc-pVTZ optimized geometry of the (R) Ibu – (R) Pro complex.	9
TS5 B97-D3/aug-cc-pVTZ optimized geometry of the (S) Nor – (S) Phe complex.	10
TS6 B97-D3/aug-cc-pVTZ optimized geometry of the (R) Nor – (R) Pro complex.	11
TS7 B97-D3/aug-cc-pVTZ optimized geometry of the (S) Bac – (R) Phe complex.	12
TS8 B97-D3/aug-cc-pVTZ optimized geometry of the (R) Bac – (S) Pro complex.	13
TS9 B97-D3/aug-cc-pVTZ optimized geometry of the (R) Ibu – (S) Phe complex.	14
TS10B97-D3/aug-cc-pVTZ optimized geometry of the (R) Ibu – (S) Pro complex.	15
TS11B97-D3/aug-cc-pVTZ optimized geometry of the (S) Nor – (R) Phe complex.	16
TS12B97-D3/aug-cc-pVTZ optimized geometry of the (S) Nor – (R) Pro complex.	17
TS13B97-D3/aug-cc-pVTZ optimized geometry of the baclofen.	18
TS14B97-D3/aug-cc-pVTZ optimized geometry of the ibuprofen.	19
TS15B97-D3/aug-cc-pVTZ optimized geometry of the norepinephrine.	20
TS16B97-D3/aug-cc-pVTZ optimized geometry of the phenethylamine	20
TS17B97-D3/aug-cc-pVTZ optimized geometry of the proline.	21
TS18Components of the stabilization energy as well as thermodynamic parameters for the complexes under study.	21
TS19IQA contributions to the “interaction” energy between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule, $E_{\text{int}}^{\text{IQA}}$, for Bac – Phe complexes and their differences (see the last column).	22
TS20IQA contributions to the “interaction” energy between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule, $E_{\text{int}}^{\text{IQA}}$, for Bac – Pro complexes and their differences (see the last column).	23
TS21IQA contributions to the “interaction” energy between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule, $E_{\text{int}}^{\text{IQA}}$, for Ibu – Phe complexes and their differences (see the last column).	24
TS22IQA contributions to the “interaction” energy between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule, $E_{\text{int}}^{\text{IQA}}$, for Ibu – Pro complexes and their differences (see the last column).	25

TS23IQA contributions to the “interaction” energy between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule, $E_{\text{int}}^{\text{IQA}}$, for Nor – Phe complexes and their differences (see the last column).	26
TS24IQA contributions to the “interaction” energy between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule, $E_{\text{int}}^{\text{IQA}}$, for Nor – Pro complexes and their differences (see the last column).	27
TS25F-SAPT/jun-cc-pVDZ interaction energy partitioning for the Bac – Phe complexes.	27
TS26F-SAPT/jun-cc-pVDZ interaction energy partitioning for the Bac – Pro complexes.	28
TS27F-SAPT/jun-cc-pVDZ interaction energy partitioning for the Ibu – Phe complexes.	28
TS28F-SAPT/jun-cc-pVDZ interaction energy partitioning for the Ibu – Pro complexes.	29
TS29F-SAPT/jun-cc-pVDZ interaction energy partitioning for the Nor – Phe complexes.	29
TS30F-SAPT/jun-cc-pVDZ interaction energy partitioning for the Nor – Pro complexes.	30
TS31Assignment of the VCD spectrum of the (R) Ibu – (R)Phe pair.	32
TS32Assignment of the VCD spectrum of the (R) Ibu – (S)Phe pair.	32
TS33Assignment of the VCD spectrum of the (S) Nor – (S)Phe pair.	32
TS34Assignment of the VCD spectrum of the (S) Nor – (R)Phe pair.	33
TS35Assignment of the VCD spectrum of the (S) Bac – (S)Phe pair.	33
TS36Assignment of the VCD spectrum of the (S) Bac – (R)Phe pair.	33
TS37Assignment of the VCD spectrum of the (R) Ibu – (R)Pro pair.	34
TS38Assignment of the VCD spectrum of the (R) Ibu – (S)Pro pair.	34
TS39Assignment of the VCD spectrum of the (R) Nor – (R)Pro pair.	34
TS40Assignment of the VCD spectrum of the (S) Nor – (R)Pro pair.	34
TS41Assignment of the VCD spectrum of the (S) Bac – (S)Pro pair.	35
TS42Assignment of the VCD spectrum of the (R) Bac – (S)Pro pair.	35

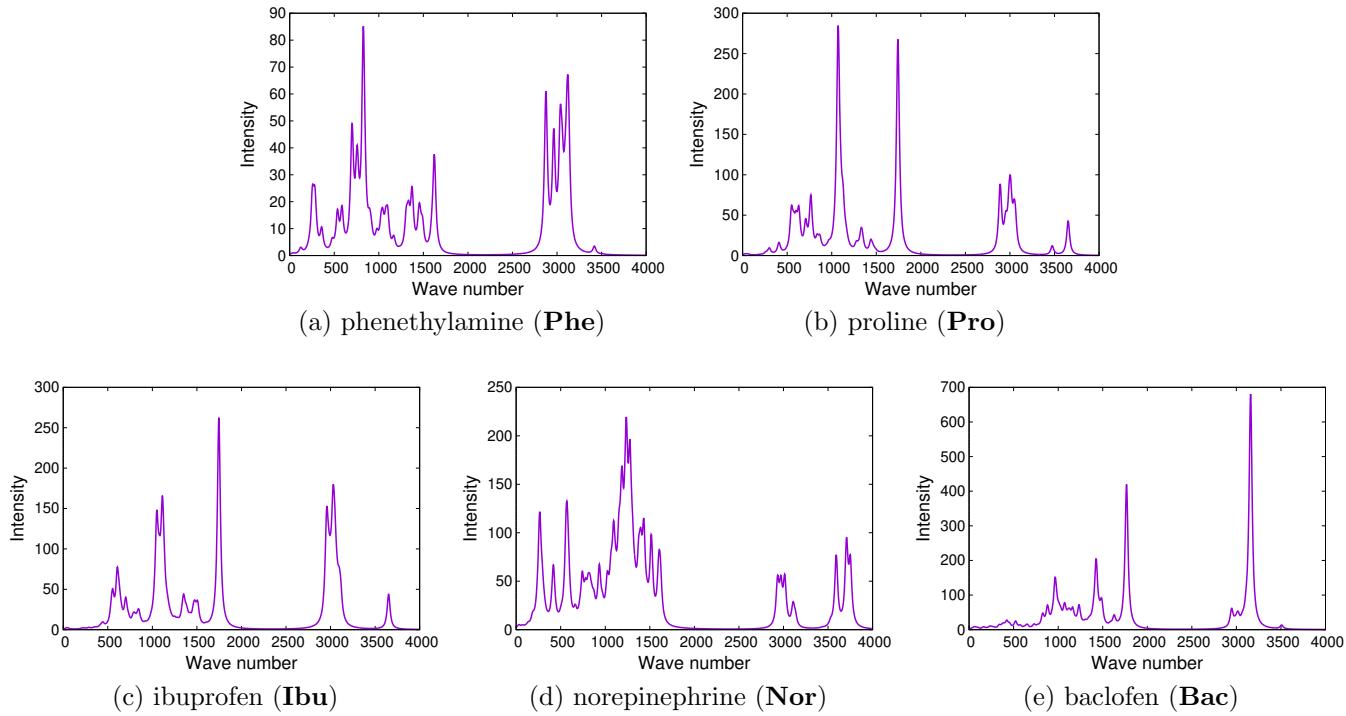


Figure FS1: Simulated B97-D3/aug-cc-pVTZ IR spectra for the monomers under study.

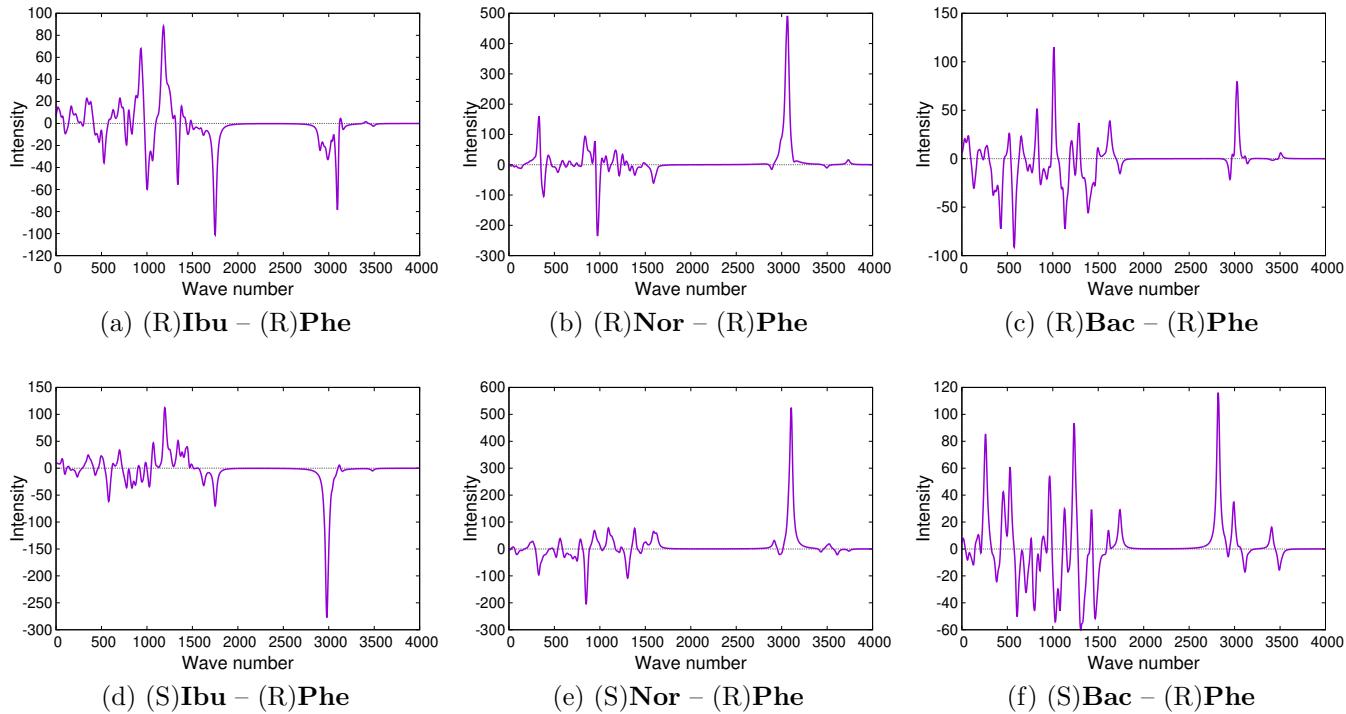


Figure FS2: Simulated B97-D3/aug-cc-pVTZ VCD spectra for complexes with phenethylamine.

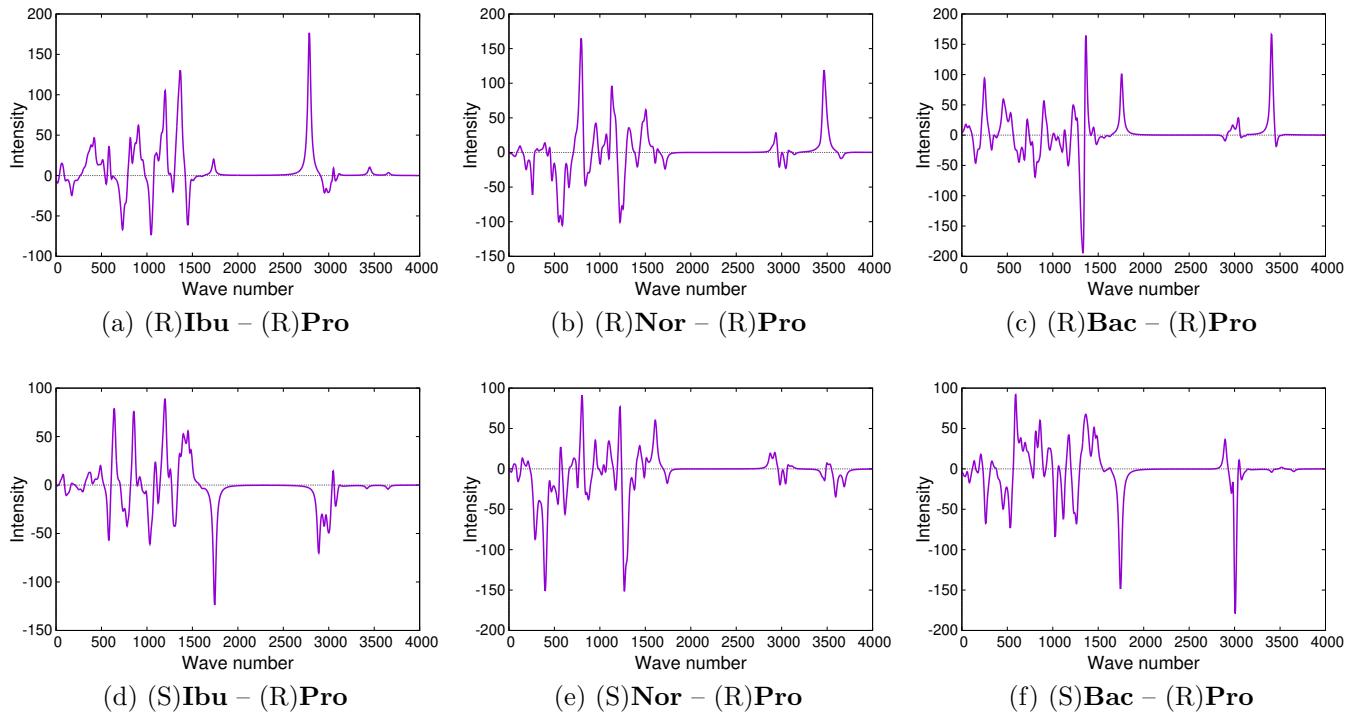


Figure FS3: Simulated B97-D3/aug-cc-pVTZ VCD spectra for complexes with proline.

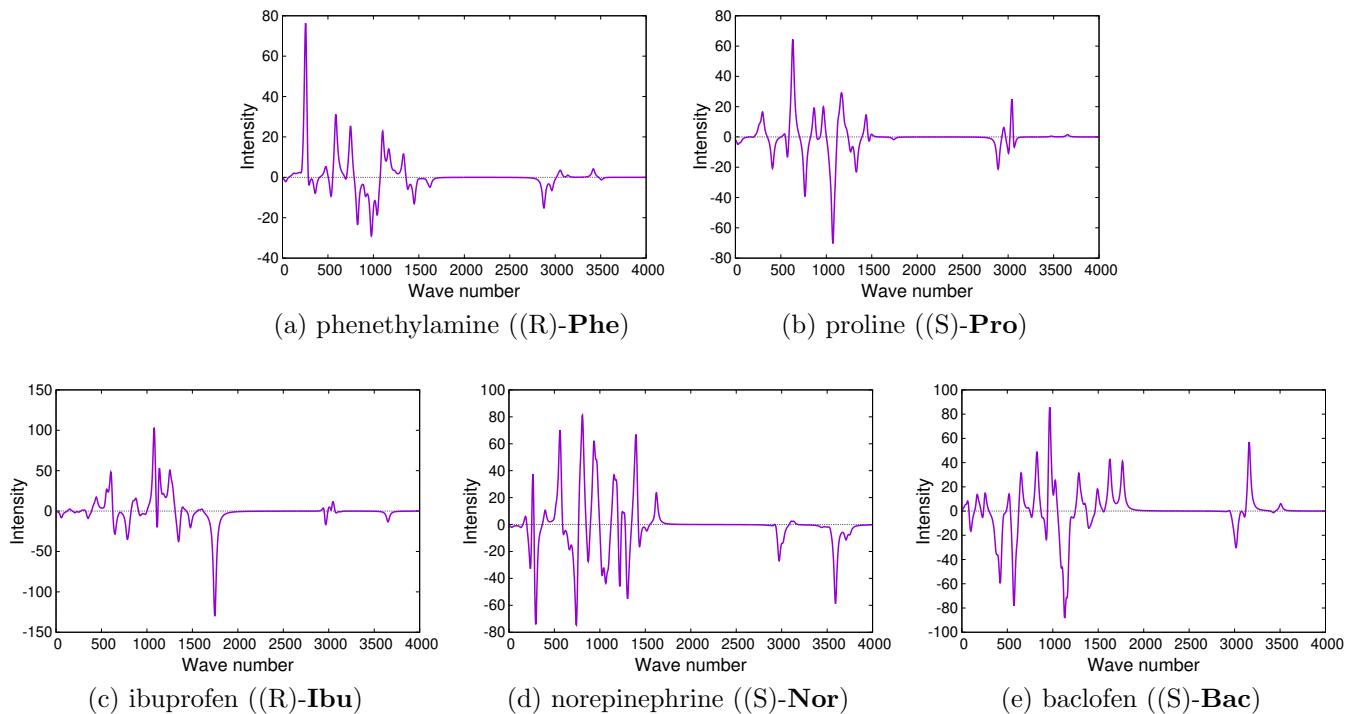


Figure FS4: Simulated B97-D3/aug-cc-pVTZ VCD spectra for the monomers under study.

Table TS1: B97-D3/aug-cc-pVTZ optimized geometry of the (S)**Bac** – (S)**Phe** complex.

At.	X	Y	Z	At.	X	Y	Z
C	5.071 623 1	-3.405 941 3	0.083 628 7	C	0.709 173 3	-0.014 141 9	-0.493 074 2
C	6.204 299 6	-3.284 759 0	1.089 627 5	C	2.230 795 4	-0.270 978 1	-0.543 494 5
C	6.513 254 3	-2.025 120 0	1.622 887 7	C	2.792 258 4	-0.419 143 5	0.863 184 6
C	7.566 567 2	-1.850 286 3	2.520 983 0	C	3.568 715 1	0.598 682 1	1.431 760 6
C	8.332 176 8	-2.955 374 3	2.893 911 1	C	4.061 679 6	0.485 958 5	2.736 010 2
C	8.048 505 5	-4.224 253 7	2.388 294 8	C	3.789 904 9	-0.658 095 4	3.488 917 7
Cl	9.660 488 0	-2.751 012 1	4.025 790 2	C	3.018 077 0	-1.681 250 2	2.930 074 6
C	6.985 399 8	-4.377 833 3	1.495 084 7	C	2.517 049 7	-1.560 515 1	1.632 369 1
C	5.594 002 9	-3.499 481 0	-1.368 929 1	N	2.603 952 2	-1.434 759 2	-1.380 818 3
N	5.962 150 8	-4.887 567 9	-1.746 810 8	H	2.284 877 9	-1.268 786 2	-2.332 732 9
H	6.320 274 7	-4.906 904 2	-2.697 824 0	H	2.099 142 4	-2.261 948 9	-1.066 016 1
H	6.709 354 6	-5.230 698 5	-1.148 890 9	H	0.315 124 2	0.133 237 9	-1.505 756 9
C	4.092 110 5	-4.544 184 4	0.477 282 9	H	0.475 309 9	0.873 751 5	0.106 432 3
C	2.948 186 4	-4.752 497 4	-0.512 840 0	H	0.198 808 8	-0.874 828 1	-0.046 038 9
O	1.808 622 7	-4.379 536 3	-0.305 075 3	H	2.711 374 9	0.604 104 0	-1.000 128 1
O	3.274 903 5	-5.392 721 7	-1.652 071 3	H	3.790 994 9	1.489 988 8	0.847 732 4
H	4.273 854 2	-5.481 777 1	-1.703 366 6	H	4.663 087 3	1.287 486 9	3.158 400 4
H	4.502 443 3	-2.472 991 9	0.112 598 2	H	4.181 352 6	-0.755 940 1	4.498 038 5
H	5.904 994 9	-1.167 002 0	1.349 933 5	H	2.805 791 8	-2.578 219 3	3.507 046 4
H	7.786 623 6	-0.871 499 9	2.935 058 4	H	1.929 450 1	-2.372 646 3	1.211 435 1
H	8.644 404 4	-5.077 101 7	2.697 192 5				
H	6.761 132 2	-5.379 531 4	1.136 564 2				
H	4.791 046 4	-3.178 015 8	-2.039 849 4				
H	6.427 757 1	-2.793 292 2	-1.485 266 6				
H	4.632 628 4	-5.491 125 3	0.576 911 3				
H	3.655 276 3	-4.298 444 9	1.446 835 8				

Table TS2: B97-D3/aug-cc-pVTZ optimized geometry of the (S)**Bac** – (S)**Pro** complex.

At.	X	Y	Z	At.	X	Y	Z
C	1.015 016 7	0.919 810 7	0.787 043 5	C	-1.766 290 3	-2.612 491 5	2.547 015 5
C	-0.355 147 4	1.114 520 9	0.962 372 7	C	-0.286 240 1	-2.187 111 0	2.766 726 2
C	-1.250 561 6	0.303 938 3	0.262 327 6	C	-0.386 115 6	-1.199 000 4	3.947 085 5
C	-0.795 385 2	-0.685 515 1	-0.606 712 2	N	-1.711 468 1	-0.545 590 6	3.835 135 3
C	0.581 369 1	-0.856 920 5	-0.774 478 8	C	-2.598 034 8	-1.682 365 2	3.484 655 5
C	1.508 176 7	-0.060 499 9	-0.087 667 6	H	-1.678 157 6	0.050 210 2	3.006 354 0
C	3.000 256 4	-0.263 911 2	-0.270 908 5	C	0.726 748 1	-0.175 361 7	4.114 411 8
C	3.653 868 5	-0.814 755 0	1.014 539 5	O	1.900 186 3	-0.365 211 7	3.858 489 6
C	3.280 816 9	-2.275 906 1	1.254 619 1	O	0.323 983 3	1.003 492 8	4.635 820 1
O	3.366 978 4	-3.120 695 0	0.389 492 9	H	-0.644 769 9	0.931 049 2	4.779 791 0
O	2.875 500 5	-2.609 156 3	2.503 675 5	H	-1.928 997 9	-3.667 065 2	2.788 756 0
C	3.691 300 9	1.035 854 7	-0.740 523 9	H	-2.054 756 1	-2.463 469 2	1.502 857 2
N	5.098 002 8	0.779 391 5	-1.078 704 0	H	0.107 791 0	-1.667 284 2	1.890 869 5
Cl	-2.980 393 0	0.518 093 9	0.511 638 7	H	0.381 025 6	-3.023 790 0	2.975 192 1
H	2.655 037 2	-1.807 339 4	3.028 007 9	H	-0.393 436 0	-1.768 004 7	4.891 303 3
H	5.161 708 8	0.188 004 0	-1.903 079 7	H	-2.865 255 4	-2.202 921 5	4.411 747 7
H	5.579 575 8	1.646 108 7	-1.296 555 5	H	-3.514 704 9	-1.309 259 4	3.022 481 1
H	1.703 915 9	1.536 780 8	1.356 545 1				
H	-0.724 185 7	1.885 437 4	1.631 867 3				
H	-1.504 052 6	-1.314 588 3	-1.135 461 6				
H	0.940 294 2	-1.637 081 2	-1.440 944 0				
H	3.149 246 1	-1.019 841 0	-1.052 014 7				
H	3.403 423 1	-0.182 474 5	1.870 848 4				
H	4.741 548 3	-0.784 933 0	0.879 953 6				
H	3.099 838 2	1.456 027 5	-1.572 158 5				
H	3.672 343 4	1.775 822 0	0.069 585 9				

Table TS3: B97-D3/aug-cc-pVTZ optimized geometry of the (R)**Ibu** – (R)**Phe** complex.

At.	X	Y	Z	At.	X	Y	Z
C	-1.499 135 9	0.332 860 5	-1.369 405 5	C	2.332 595 6	-0.516 009 0	-6.280 534 9
C	-0.240 749 4	0.250 432 4	-2.252 415 3	C	2.466 547 2	-1.169 009 0	-7.515 464 5
C	0.034 386 7	-1.187 572 1	-2.683 109 8	C	2.915 554 3	-2.488 233 0	-7.584 648 2
C	-0.651 925 3	-1.726 098 0	-3.779 059 9	C	3.248 853 0	-3.174 739 4	-6.412 641 5
C	-0.430 650 8	-3.039 447 3	-4.194 141 1	C	3.125 373 5	-2.530 968 5	-5.181 053 6
C	0.480 708 7	-3.863 214 3	-3.523 131 9	C	2.666 226 1	-1.212 383 7	-5.112 976 4
C	1.141 448 0	-3.332 827 7	-2.406 610 6	C	1.787 909 5	0.904 777 0	-6.264 727 4
C	0.925 363 2	-2.018 980 2	-1.990 133 1	C	0.256 420 9	0.918 962 1	-6.384 386 1
C	0.745 705 5	-5.281 041 9	-3.968 414 2	N	2.176 505 5	1.625 692 7	-5.027 046 1
C	0.081 952 2	-6.357 446 8	-3.074 548 3	H	3.191 624 3	1.643 146 3	-4.945 320 2
C	-1.448 116 6	-6.246 767 8	-3.103 932 4	H	1.877 170 6	2.595 471 1	-5.098 085 9
C	0.540 204 5	-7.758 481 5	-3.502 295 3	H	2.217 061 3	-0.636 314 8	-8.431 807 5
C	0.976 943 1	0.840 703 6	-1.518 778 9	H	3.012 583 1	-2.976 788 2	-8.551 101 6
O	1.064 899 9	0.879 177 5	-0.311 015 3	H	3.601 747 6	-4.201 704 9	-6.461 578 4
O	1.989 025 8	1.284 173 0	-2.294 042 6	H	3.367 595 2	-3.055 765 8	-4.262 059 5
H	1.811 047 9	1.244 319 8	-3.276 584 3	H	2.554 386 9	-0.742 863 7	-4.142 179 8
H	-2.355 582 0	-0.093 212 9	-1.901 816 8	H	2.197 893 4	1.418 611 1	-7.150 128 7
H	-1.722 268 1	1.371 801 9	-1.105 309 9	H	-0.186 417 8	0.395 318 3	-5.533 862 2
H	-1.340 993 2	-0.227 177 3	-0.444 509 3	H	-0.054 893 9	0.413 325 2	-7.303 055 5
H	-0.405 794 7	0.844 349 6	-3.162 012 1	H	-0.124 630 2	1.947 635 0	-6.409 032 1
H	-1.369 181 3	-1.109 526 3	-4.316 694 7				
H	-0.961 352 7	-3.426 217 4	-5.061 663 6				
H	1.851 933 3	-3.953 859 4	-1.863 663 3				
H	1.460 152 0	-1.634 550 1	-1.125 347 0				
H	0.396 133 5	-5.418 686 5	-5.000 742 8				
H	1.830 049 4	-5.457 476 3	-3.978 185 9				
H	0.419 533 6	-6.180 791 8	-2.042 867 6				
H	-1.822 574 3	-6.413 251 7	-4.122 927 9				
H	-1.784 115 0	-5.256 830 2	-2.779 076 3				
H	-1.907 056 9	-6.997 654 9	-2.450 661 2				
H	0.234 177 5	-7.962 708 3	-4.536 763 3				
H	0.096 538 9	-8.531 051 0	-2.863 673 8				
H	1.631 186 1	-7.856 104 8	-3.448 776 1				

Table TS4: B97-D3/aug-cc-pVTZ optimized geometry of the (R)**Ibu** – (R)**Pro** complex.

At.	X	Y	Z	At.	X	Y	Z
C	-1.522 857 8	0.503 710 9	-1.099 841 2	C	3.720 964 3	-0.479 471 4	-6.219 425 9
C	-0.337 278 4	0.373 083 2	-2.070 805 8	C	4.728 567 7	-0.336 358 5	-5.069 008 3
C	-0.145 548 2	-1.059 679 1	-2.528 434 4	C	4.037 201 5	-1.112 181 8	-3.915 115 3
C	-0.591 635 0	-1.482 277 3	-3.784 940 6	N	2.589 166 6	-0.916 951 5	-4.133 227 8
C	-0.373 423 5	-2.791 579 5	-4.223 691 0	C	2.377 596 1	-0.335 038 5	-5.491 013 7
C	0.299 801 4	-3.720 217 8	-3.419 631 9	H	2.107 426 5	-1.808 788 9	-4.055 549 9
C	0.719 545 9	-3.300 357 7	-2.146 738 0	C	4.420 562 1	-2.589 365 5	-3.940 009 6
C	0.495 739 1	-1.997 808 9	-1.705 753 3	O	3.709 000 8	-3.501 206 8	-4.302 229 1
C	0.555 356 3	-5.134 826 2	-3.878 420 3	O	5.697 162 1	-2.769 149 1	-3.499 051 4
C	-0.379 605 1	-6.179 632 0	-3.220 908 0	H	5.882 294 6	-3.722 057 8	-3.551 692 9
C	-1.847 347 5	-5.937 692 9	-3.597 510 0	H	3.863 236 0	0.268 975 1	-7.004 435 5
C	0.059 050 2	-7.599 576 5	-3.604 037 2	H	3.801 602 0	-1.474 989 1	-6.671 025 6
C	0.944 210 3	1.001 093 6	-1.463 721 1	H	4.827 348 4	0.713 831 0	-4.774 076 3
O	0.920 080 8	1.680 873 2	-0.459 717 7	H	5.725 031 7	-0.723 633 3	-5.292 069 2
O	2.092 993 7	0.806 920 0	-2.135 413 7	H	4.340 204 7	-0.711 045 0	-2.941 077 4
H	2.062 533 3	0.130 370 9	-2.894 582 5	H	1.549 696 3	-0.844 034 1	-5.991 440 6
H	-2.438 498 7	0.141 307 3	-1.578 830 9	H	2.111 122 8	0.724 756 3	-5.384 242 2
H	-1.655 432 0	1.544 129 4	-0.793 489 7				
H	-1.341 282 4	-0.092 309 6	-0.200 242 1				
H	-0.548 624 7	0.982 633 9	-2.962 252 6				
H	-1.093 534 0	-0.772 478 7	-4.440 232 4				
H	-0.716 729 8	-3.093 719 7	-5.211 025 9				
H	1.246 556 3	-4.002 859 0	-1.504 088 0				
H	0.856 027 4	-1.690 300 0	-0.726 286 1				
H	0.440 665 5	-5.200 052 0	-4.968 904 3				
H	1.596 685 3	-5.397 809 2	-3.655 763 8				
H	-0.281 662 6	-6.068 326 5	-2.131 063 4				
H	-1.983 626 7	-6.039 631 7	-4.682 547 6				
H	-2.174 949 3	-4.933 264 6	-3.310 248 0				
H	-2.503 088 8	-6.665 981 7	-3.106 845 7				
H	-0.009 554 5	-7.741 717 2	-4.690 478 6				
H	-0.579 945 5	-8.351 783 8	-3.127 155 9				
H	1.096 194 3	-7.791 531 8	-3.304 707 1				

Table TS5: B97-D3/aug-cc-pVTZ optimized geometry of the (S)**Nor** – (S)**Phe** complex.

At.	X	Y	Z	At.	X	Y	Z
C	6.330 080 2	-3.497 409 1	1.620 417 0	C	0.958 676 0	-0.046 215 1	-0.195 358 5
C	5.859 725 8	-2.621 728 9	0.638 534 2	C	2.479 387 5	-0.110 262 3	0.020 603 8
C	6.248 607 5	-1.277 016 2	0.607 752 6	C	2.868 980 9	-0.574 870 2	1.415 146 1
C	7.165 579 3	-0.827 087 6	1.587 058 4	C	3.780 951 8	0.165 206 0	2.175 629 0
C	7.639 116 9	-1.710 633 5	2.556 637 0	C	4.195 238 0	-0.285 039 5	3.430 667 3
C	7.223 648 4	-3.049 840 0	2.598 413 5	C	3.700 673 2	-1.485 284 1	3.944 271 1
C	7.669 150 5	-3.944 579 3	3.734 778 0	C	2.787 535 9	-2.231 824 7	3.194 328 7
O	7.828 563 2	-5.320 916 3	3.337 366 6	C	2.375 223 1	-1.777 861 2	1.939 741 9
H	8.278 173 3	-5.326 415 3	2.482 855 2	N	3.173 129 1	-0.976 194 0	-0.981 282 3
C	6.651 430 2	-3.956 271 1	4.904 918 4	H	2.882 301 7	-0.719 919 7	-1.922 809 0
N	5.320 058 8	-4.473 582 4	4.591 023 6	H	2.892 600 2	-1.944 465 0	-0.840 937 1
H	4.907 842 2	-3.916 051 7	3.846 447 7	H	0.731 311 7	0.338 135 9	-1.196 920 0
H	5.418 325 1	-5.415 162 2	4.219 765 7	H	0.490 555 4	0.613 824 6	0.542 941 6
O	7.535 916 5	0.498 023 2	1.541 689 6	H	0.504 619 9	-1.038 729 9	-0.100 978 6
H	8.181 069 9	0.659 672 0	2.241 830 9	H	2.892 180 7	0.894 967 9	-0.125 293 6
O	5.791 520 8	-0.389 461 5	-0.309 967 9	H	4.203 519 0	1.078 172 0	1.764 638 1
H	4.909 327 4	-0.701 839 8	-0.661 906 9	H	4.918 992 0	0.294 870 2	3.996 977 5
H	6.004 682 0	-4.533 114 1	1.626 544 6	H	4.029 509 8	-1.849 971 6	4.913 386 9
H	5.169 889 4	-2.975 321 9	-0.122 804 0	H	2.403 349 7	-3.169 423 1	3.588 007 0
H	8.336 630 3	-1.340 571 5	3.309 621 7	H	1.665 312 2	-2.372 847 5	1.368 639 8
H	8.625 052 0	-3.562 331 6	4.135 339 9				
H	6.556 976 0	-2.928 838 5	5.277 611 4				
H	7.078 586 2	-4.564 791 3	5.711 673 4				

Table TS6: B97-D3/aug-cc-pVTZ optimized geometry of the (R)**Nor** – (R)**Pro** complex.

At.	X	Y	Z	At.	X	Y	Z
C	-0.286 697 8	-2.765 713 0	-2.207 864 1	C	1.418 384 5	-6.187 890 1	-0.450 588 7
C	0.170 882 9	-3.449 656 4	-3.338 782 6	C	2.484 134 8	-5.350 392 3	-1.203 085 7
C	-0.324 234 2	-4.717 888 1	-3.634 810 8	C	3.015 651 9	-6.344 291 0	-2.293 916 0
C	-1.304 569 0	-5.297 390 5	-2.804 584 6	N	1.915 574 9	-7.254 181 7	-2.603 875 2
C	-1.756 756 6	-4.611 525 4	-1.679 532 6	C	1.237 420 3	-7.478 234 2	-1.310 989 5
C	-1.246 830 5	-3.343 985 4	-1.368 969 9	H	1.275 769 6	-6.795 707 4	-3.246 179 3
C	-1.660 584 2	-2.647 858 1	-0.090 714 0	C	3.520 221 9	-5.593 333 2	-3.503 687 6
C	-0.833 952 4	-3.159 548 5	1.111 488 8	O	2.866 075 3	-5.268 186 3	-4.480 656 4
N	0.582 491 2	-2.801 873 7	0.961 682 3	O	4.823 546 6	-5.246 419 2	-3.370 210 2
O	-3.060 526 1	-2.896 830 2	0.132 834 3	H	5.059 782 9	-4.717 375 4	-4.151 893 0
O	-1.796 112 3	-6.547 645 1	-3.077 079 5	H	1.744 724 2	-6.433 512 0	0.565 366 3
O	0.083 575 8	-5.470 949 7	-4.717 217 7	H	0.480 384 9	-5.633 179 5	-0.377 517 7
H	0.976 167 9	-3.292 904 1	0.163 393 7	H	2.016 986 4	-4.492 678 1	-1.695 310 8
H	1.108 293 9	-3.099 653 0	1.777 880 9	H	3.289 988 6	-4.985 681 0	-0.559 769 9
H	-3.306 169 8	-2.475 708 8	0.966 073 2	H	3.850 607 2	-6.918 117 4	-1.872 688 3
H	-1.374 385 0	-6.852 778 9	-3.895 705 4	H	1.711 598 0	-8.332 428 5	-0.811 475 0
H	1.041 522 1	-5.327 041 6	-4.856 202 6	H	0.187 865 1	-7.727 755 5	-1.485 308 4
H	0.108 307 4	-1.780 422 0	-1.976 416 2				
H	0.918 503 3	-3.009 988 3	-3.993 696 4				
H	-2.507 338 7	-5.072 145 1	-1.045 445 1				
H	-1.471 204 9	-1.568 465 3	-0.197 063 8				
H	-1.028 785 5	-4.241 745 4	1.208 060 8				
H	-1.211 314 3	-2.671 921 5	2.021 456 9				

Table TS7: B97-D3/aug-cc-pVTZ optimized geometry of the (S)**Bac** – (R)**Phe** complex.

At.	X	Y	Z	At.	X	Y	Z
C	0.894 243 2	1.644 457 5	0.015 252 5	C	-0.795 532 2	-1.400 017 5	3.543 334 7
C	-0.458 335 1	1.863 362 7	0.285 513 0	C	-2.179 359 2	-1.291 870 1	3.360 289 5
C	-1.316 363 2	0.767 058 8	0.391 811 3	C	-2.896 625 4	-2.290 195 9	2.695 029 5
C	-0.838 023 2	-0.533 500 8	0.234 774 2	C	-2.231 657 0	-3.412 625 6	2.197 704 0
C	0.516 041 4	-0.728 091 3	-0.046 478 5	C	-0.848 540 4	-3.529 090 1	2.373 597 8
C	1.406 297 4	0.348 889 4	-0.162 325 4	C	-0.136 237 5	-2.533 913 9	3.043 796 2
C	2.887 016 9	0.102 210 5	-0.386 796 0	C	-0.040 428 0	-0.298 287 9	4.270 746 8
C	3.637 714 9	0.024 785 4	0.970 311 1	C	0.692 315 7	-0.807 611 9	5.518 833 5
C	3.528 791 8	-1.363 831 5	1.603 241 7	N	0.958 374 6	0.340 278 6	3.356 342 6
O	4.001 211 9	-2.341 404 1	1.058 670 7	H	0.505 208 0	0.616 570 6	2.487 273 1
O	2.917 284 2	-1.458 741 4	2.798 200 1	H	1.306 527 6	1.193 720 0	3.788 169 3
C	3.531 020 5	1.140 125 1	-1.329 193 3	H	-2.702 463 0	-0.411 183 5	3.726 198 2
N	4.912 354 1	0.756 536 0	-1.645 607 4	H	-3.969 037 6	-2.182 651 0	2.555 716 5
Cl	-3.017 835 5	1.038 264 3	0.736 093 1	H	-2.784 455 7	-4.190 138 8	1.676 278 8
H	2.287 692 7	-0.685 176 2	2.987 430 8	H	-0.320 456 5	-4.397 175 8	1.987 003 7
H	4.932 515 5	-0.131 913 7	-2.139 100 7	H	0.937 760 8	-2.633 528 0	3.157 772 6
H	5.349 327 3	1.446 250 4	-2.248 446 9	H	-0.782 428 8	0.450 421 3	4.587 386 1
H	1.551 673 0	2.506 331 5	-0.052 927 2	H	1.436 186 6	-1.562 318 1	5.247 939 3
H	-0.845 322 5	2.869 582 7	0.412 294 8	H	-0.020 303 5	-1.252 785 4	6.219 831 4
H	-1.505 939 1	-1.378 590 0	0.356 709 8	H	1.207 276 6	0.017 118 1	6.026 677 9
H	0.888 744 1	-1.743 820 3	-0.152 523 4				
H	3.003 436 8	-0.879 231 8	-0.864 911 4				
H	3.281 677 0	0.814 148 3	1.640 884 9				
H	4.703 264 9	0.183 560 3	0.772 438 0				
H	2.878 907 9	1.248 114 5	-2.213 162 4				
H	3.570 477 5	2.120 909 0	-0.838 075 1				

Table TS8: B97-D3/aug-cc-pVTZ optimized geometry of the (R)**Bac** – (S)**Pro** complex.

At.	X	Y	Z	At.	X	Y	Z
C	-0.473 994 8	0.769 112 9	0.411 223 2	C	1.002 142 1	-0.345 922 2	3.869 770 7
C	0.917 064 0	0.661 856 5	0.346 075 4	C	1.609 599 1	-1.569 326 7	3.134 695 4
C	1.484 892 4	-0.387 274 8	-0.375 972 7	C	0.372 486 9	-2.298 010 5	2.536 935 3
C	0.682 193 8	-1.315 855 6	-1.039 339 2	N	-0.757 689 1	-1.992 171 9	3.438 716 0
C	-0.706 050 0	-1.186 837 1	-0.968 907 9	C	-0.535 118 7	-0.589 456 4	3.860 136 4
C	-1.310 708 2	-0.149 043 3	-0.241 186 7	H	-0.683 455 3	-2.616 940 3	4.241 561 0
C	-2.824 173 6	-0.093 630 2	-0.124 710 5	C	0.587 170 2	-3.795 610 2	2.448 888 0
C	-3.304 985 9	-0.954 769 6	1.067 691 8	O	0.359 823 9	-4.572 993 1	3.350 515 9
C	-3.143 964 8	-2.450 129 9	0.792 769 4	O	1.121 535 7	-4.157 634 8	1.255 025 9
O	-3.552 528 5	-2.959 099 5	-0.231 425 2	H	1.256 522 6	-5.120 398 7	1.287 724 3
O	-2.549 425 7	-3.189 918 4	1.746 962 9	H	1.248 228 3	0.588 546 1	3.357 211 1
C	-3.381 138 7	1.340 046 2	-0.033 894 4	H	1.387 663 1	-0.274 071 0	4.890 942 0
N	-4.847 281 9	1.324 591 9	-0.130 874 3	H	2.323 889 1	-1.293 119 6	2.356 351 4
Cl	3.233 942 8	-0.551 115 6	-0.429 708 2	H	2.114 070 9	-2.233 108 9	3.844 617 7
H	-2.094 938 0	-2.628 125 7	2.444 673 9	H	0.147 135 1	-1.899 557 4	1.545 235 8
H	-5.137 650 2	0.999 278 9	-1.049 041 8	H	-1.013 691 8	-0.411 299 3	4.827 017 4
H	-5.225 844 7	2.259 081 0	-0.012 693 7	H	-1.007 807 9	0.067 249 1	3.122 797 2
H	-0.900 497 3	1.587 729 9	0.983 628 3				
H	1.555 276 3	1.379 047 5	0.852 166 4				
H	1.137 097 9	-2.133 469 7	-1.588 124 6				
H	-1.334 659 0	-1.920 350 3	-1.466 422 8				
H	-3.246 683 1	-0.554 738 3	-1.026 676 1				
H	-4.376 844 5	-0.771 340 2	1.207 611 2				
H	-2.788 463 7	-0.651 175 5	1.983 309 9				
H	-3.124 411 1	1.781 943 9	0.937 872 7				
H	-2.886 728 2	1.953 010 3	-0.807 148 8				

Table TS9: B97-D3/aug-cc-pVTZ optimized geometry of the (R)**Ibu** – (S)**Phe** complex.

At.	X	Y	Z	At.	X	Y	Z
C	-0.883 623 3	0.098 625 8	-0.538 108 5	C	2.218 184 8	-0.700 473 4	-5.743 563 5
C	0.017 207 0	0.294 184 5	-1.770 012 7	C	3.368 023 6	-0.524 002 4	-4.965 558 1
C	0.128 893 0	-0.996 014 1	-2.572 583 1	C	4.327 055 1	-1.538 142 8	-4.870 265 1
C	-0.862 849 0	-1.326 706 9	-3.504 631 3	C	4.153 016 8	-2.742 040 7	-5.553 197 9
C	-0.815 272 7	-2.530 789 3	-4.210 095 4	C	3.006 409 8	-2.929 429 3	-6.330 783 8
C	0.225 944 4	-3.444 921 2	-4.006 628 4	C	2.049 363 6	-1.920 422 4	-6.417 166 0
C	1.214 218 2	-3.112 728 7	-3.070 599 5	C	1.138 786 2	0.362 888 8	-5.893 169 7
C	1.168 766 8	-1.912 067 0	-2.363 933 8	C	1.015 956 9	0.839 696 6	-7.355 064 6
C	0.282 685 7	-4.762 855 5	-4.739 077 2	N	1.342 403 7	1.502 953 6	-4.964 370 1
C	-0.184 763 4	-5.970 526 9	-3.888 940 5	H	0.547 122 8	2.132 540 5	-5.042 760 7
C	-1.663 796 7	-5.848 510 6	-3.499 535 3	H	2.150 036 9	2.045 760 8	-5.269 423 9
C	0.079 595 7	-7.283 262 2	-4.638 966 6	H	3.519 388 8	0.386 883 4	-4.397 111 5
C	1.404 568 6	0.823 576 2	-1.354 800 6	H	5.206 777 1	-1.383 777 7	-4.250 711 5
O	1.815 799 2	0.786 560 4	-0.215 328 8	H	4.898 605 3	-3.529 378 0	-5.476 951 5
O	2.160 481 7	1.326 518 3	-2.349 452 9	H	2.850 822 9	-3.866 551 1	-6.858 991 8
H	1.721 074 8	1.271 998 0	-3.252 424 4	H	1.147 457 3	-2.090 956 8	-7.000 094 9
H	-1.872 956 1	-0.250 120 4	-0.851 675 8	H	0.191 231 8	-0.100 797 0	-5.592 854 0
H	-0.989 502 7	1.035 133 4	0.018 023 2	H	1.951 152 8	1.314 980 4	-7.674 880 4
H	-0.442 419 0	-0.644 095 6	0.131 533 2	H	0.201 407 7	1.567 695 2	-7.452 615 0
H	-0.434 850 5	1.054 551 9	-2.423 240 1	H	0.812 960 0	0.005 646 4	-8.033 504 0
H	-1.684 041 9	-0.632 853 4	-3.679 488 3				
H	-1.595 926 6	-2.763 444 7	-4.932 094 6				
H	2.043 996 2	-3.796 898 9	-2.907 477 1				
H	1.955 514 9	-1.679 911 7	-1.650 474 9				
H	-0.334 981 4	-4.715 760 5	-5.647 125 3				
H	1.314 576 2	-4.944 250 1	-5.066 008 4				
H	0.414 428 5	-5.970 795 3	-2.966 757 7				
H	-2.295 367 4	-5.840 831 9	-4.398 223 5				
H	-1.856 621 4	-4.925 079 7	-2.944 190 3				
H	-1.975 577 7	-6.695 294 7	-2.877 494 4				
H	-0.489 609 8	-7.311 526 9	-5.577 566 7				
H	-0.224 009 5	-8.148 493 2	-4.038 296 1				
H	1.142 085 2	-7.395 220 7	-4.886 205 1				

Table TS10: B97-D3/aug-cc-pVTZ optimized geometry of the (R)**Ibu** – (S)**Pro** complex.

At.	X	Y	Z	At.	X	Y	Z
C	-1.328 913 4	0.354 191 7	-1.150 160 7	C	1.252 049 5	-0.377 370 1	-6.725 109 4
C	-0.095 247 2	0.338 737 1	-2.068 758 9	C	2.571 036 1	-1.124 033 9	-6.464 677 0
C	0.146 363 0	-1.053 396 5	-2.633 893 3	C	2.825 909 9	-0.824 017 3	-4.965 851 7
C	-0.525 214 5	-1.468 309 1	-3.790 251 1	N	2.373 664 9	0.568 320 3	-4.761 691 4
C	-0.336 943 2	-2.749 021 7	-4.314 243 7	C	1.292 178 3	0.835 379 5	-5.756 321 5
C	0.532 475 1	-3.659 772 8	-3.700 206 3	H	3.172 641 7	1.171 528 1	-4.950 979 2
C	1.189 896 3	-3.248 235 3	-2.531 188 1	C	4.290 674 7	-0.968 893 1	-4.608 111 0
C	0.999 221 9	-1.970 898 4	-2.003 097 7	O	5.105 870 5	-0.072 764 4	-4.643 521 3
C	0.736 590 5	-5.052 984 2	-4.245 178 8	O	4.600 722 8	-2.254 975 7	-4.291 435 7
C	-0.012 773 9	-6.150 389 8	-3.447 926 6	H	5.557 724 2	-2.276 586 8	-4.118 703 0
C	-1.532 423 6	-5.943 976 5	-3.495 781 9	H	1.148 412 7	-0.068 636 2	-7.769 295 0
C	0.369 687 3	-7.541 183 4	-3.971 918 3	H	0.406 853 5	-1.024 888 8	-6.476 585 1
C	1.149 009 9	0.877 902 8	-1.329 564 7	H	3.385 375 1	-0.697 720 4	-7.063 112 9
O	1.172 037 8	1.063 323 1	-0.131 319 3	H	2.510 852 1	-2.196 309 9	-6.667 644 6
O	2.226 645 8	1.125 920 1	-2.092 365 9	H	2.231 486 8	-1.501 300 3	-4.348 880 5
H	2.110 829 5	0.893 290 3	-3.069 090 5	H	0.336 735 2	0.948 553 4	-5.234 620 4
H	-2.208 217 7	0.000 366 4	-1.698 675 2	H	1.500 668 5	1.775 110 3	-6.277 461 2
H	-1.518 320 2	1.364 085 1	-0.775 105 0				
H	-1.161 864 7	-0.298 720 4	-0.289 419 4				
H	-0.275 927 7	1.015 426 1	-2.916 116 7				
H	-1.200 100 9	-0.776 942 6	-4.291 199 3				
H	-0.867 080 0	-3.044 231 2	-5.217 748 7				
H	1.873 908 9	-3.934 663 0	-2.036 122 8				
H	1.533 995 5	-1.677 025 4	-1.103 031 0				
H	0.406 220 0	-5.097 512 3	-5.292 290 5				
H	1.808 840 4	-5.291 041 9	-4.239 571 3				
H	0.311 825 5	-6.071 974 8	-2.400 129 6				
H	-1.893 881 0	-6.005 735 5	-4.531 053 1				
H	-1.817 529 3	-4.965 151 7	-3.096 971 6				
H	-2.049 061 7	-6.715 293 2	-2.913 515 3				
H	0.070 773 2	-7.651 155 5	-5.022 703 0				
H	-0.131 202 9	-8.329 245 3	-3.398 037 4				
H	1.451 525 1	-7.708 777 0	-3.910 363 9				

Table TS11: B97-D3/aug-cc-pVTZ optimized geometry of the (S)**Nor** – (R)**Phe** complex.

At.	X	Y	Z	At.	X	Y	Z
C	-7.633 501 9	-1.578 255 8	2.526 283 4	C	-1.130 873 9	0.309 342 6	-0.300 381 0
C	-7.126 315 7	-0.615 438 0	1.647 225 1	C	-2.579 581 8	-0.200 584 2	-0.153 912 2
C	-6.312 637 1	-0.981 518 6	0.577 177 3	C	-2.858 766 8	-0.631 623 0	1.280 094 6
C	-5.988 627 5	-2.340 858 2	0.371 803 7	C	-2.958 584 4	0.359 516 9	2.269 170 2
C	-6.515 941 9	-3.288 003 7	1.252 111 5	C	-3.195 430 5	0.026 429 3	3.601 994 4
C	-7.329 390 0	-2.926 390 2	2.333 594 1	C	-3.336 484 8	-1.315 189 9	3.972 026 4
C	-7.782 186 9	-4.000 348 1	3.320 330 5	C	-3.247 019 8	-2.308 366 1	2.996 783 7
O	-8.797 913 6	-3.551 310 0	4.216 132 3	C	-3.008 311 2	-1.968 926 9	1.660 930 4
H	-8.316 055 9	-3.130 260 2	4.948 301 5	N	-2.903 529 1	-1.265 644 6	-1.130 984 7
C	-6.575 196 8	-4.521 633 9	4.135 159 5	H	-2.873 516 0	-0.877 244 1	-2.069 766 7
N	-6.109 072 5	-3.438 251 2	5.016 423 1	H	-2.191 031 1	-1.992 621 9	-1.103 127 3
H	-5.427 279 7	-3.777 503 5	5.685 981 4	H	-0.952 829 5	0.672 049 1	-1.319 760 0
H	-5.665 736 5	-2.708 604 8	4.464 237 4	H	-0.424 331 7	-0.502 346 3	-0.088 277 7
O	-5.198 149 5	-2.759 655 3	-0.653 166 8	H	-0.924 965 1	1.125 737 6	0.399 678 3
H	-4.498 682 6	-2.081 733 4	-0.870 602 6	H	-3.268 371 7	0.621 439 2	-0.379 062 8
O	-5.793 894 4	-0.062 288 0	-0.319 238 5	H	-2.859 601 3	1.406 553 7	1.987 727 5
H	-6.128 437 0	0.810 876 4	-0.080 206 8	H	-3.277 659 6	0.810 497 6	4.350 461 7
H	-8.275 107 7	-1.282 441 8	3.349 188 7	H	-3.522 174 7	-1.577 982 9	5.010 380 4
H	-7.369 513 3	0.437 010 6	1.793 130 7	H	-3.380 674 3	-3.352 559 6	3.267 199 3
H	-6.253 916 0	-4.329 653 7	1.077 445 1	H	-2.987 736 9	-2.757 613 3	0.916 365 6
H	-8.207 940 0	-4.843 411 2	2.759 527 2				
H	-5.806 112 0	-4.901 425 7	3.443 539 3				
H	-6.915 917 5	-5.352 764 9	4.763 082 9				

Table TS12: B97-D3/aug-cc-pVTZ optimized geometry of the (S)**Nor** – (R)**Pro** complex.

At.	X	Y	Z	At.	X	Y	Z
C	0.948 238 7	-2.454 625 6	3.633 227 7	C	1.163 855 1	-0.854 196 0	-0.298 075 8
C	1.412 243 6	-1.136 759 9	3.607 535 9	C	1.438 953 8	0.588 898 1	0.149 568 0
C	0.514 936 0	-0.068 463 4	3.637 660 0	C	0.064 867 0	1.274 716 8	-0.114 934 4
C	-0.872 394 2	-0.328 272 9	3.668 291 8	N	-0.944 463 6	0.221 024 2	-0.019 962 3
C	-1.318 403 3	-1.648 314 1	3.710 969 7	C	-0.274 876 4	-1.086 507 6	0.183 467 9
C	-0.421 553 0	-2.724 937 9	3.693 001 6	C	-0.169 419 4	2.446 213 8	0.829 776 1
C	-0.963 561 9	-4.151 221 2	3.661 913 7	O	-0.996 039 2	2.482 706 9	1.719 570 9
O	-0.003 849 0	-5.124 790 7	4.063 912 9	O	0.623 339 8	3.528 377 0	0.631 498 5
H	0.451 195 0	-5.380 842 9	3.242 617 3	H	-1.619 354 5	0.424 049 9	0.708 453 1
C	-1.490 220 2	-4.501 416 0	2.248 726 6	H	1.237 855 0	3.361 531 5	-0.099 472 3
N	-0.336 581 0	-4.652 505 0	1.345 990 6	H	1.221 430 3	-0.932 053 0	-1.390 298 2
H	-0.607 438 5	-5.083 578 4	0.469 361 4	H	1.868 183 8	-1.567 580 9	0.139 018 1
H	0.067 391 8	-3.744 866 7	1.135 568 8	H	1.666 681 4	0.617 701 5	1.218 698 8
O	-1.790 721 7	0.700 336 5	3.715 586 0	H	2.259 002 8	1.066 262 3	-0.396 916 3
H	-1.610 196 1	1.336 096 9	2.994 554 0	H	0.068 233 5	1.700 777 5	-1.135 016 7
O	1.020 868 8	1.209 309 0	3.627 003 4	H	-0.273 019 8	-1.382 724 9	1.239 640 4
H	0.296 495 9	1.838 379 4	3.759 076 4	H	-0.785 479 4	-1.866 240 5	-0.394 708 0
H	1.654 856 3	-3.278 131 2	3.637 765 4				
H	2.476 481 0	-0.917 288 2	3.588 437 4				
H	-2.392 627 9	-1.817 525 5	3.748 524 5				
H	-1.804 259 0	-4.220 585 1	4.365 412 4				
H	-2.006 656 3	-5.466 380 7	2.301 878 0				
H	-2.214 994 8	-3.735 570 9	1.930 675 9				

Table TS13: B97-D3/aug-cc-pVTZ optimized geometry of the baclofen.

At.	X	Y	Z
C	5.034 417 7	-3.402 037 8	0.108 244 5
C	6.179 237 8	-3.284 525 8	1.105 606 9
C	6.537 773 9	-2.022 331 3	1.602 042 5
C	7.611 514 6	-1.855 488 4	2.477 638 5
C	8.350 934 2	-2.973 428 2	2.865 726 6
C	8.020 802 4	-4.243 990 9	2.393 964 5
Cl	9.703 364 9	-2.781 087 3	3.966 327 7
C	6.938 699 2	-4.388 751 4	1.522 877 3
C	5.568 456 0	-3.471 743 2	-1.345 717 2
N	6.021 770 4	-4.836 356 3	-1.700 224 4
H	6.377 798 9	-4.850 515 8	-2.651 939 1
H	6.789 919 8	-5.120 504 0	-1.098 051 9
C	4.065 890 2	-4.557 808 4	0.474 962 5
C	2.943 437 6	-4.776 541 6	-0.543 723 9
O	1.804 563 1	-4.407 182 9	-0.370 489 6
O	3.310 868 6	-5.435 508 2	-1.670 723 0
H	4.302 757 9	-5.524 334 7	-1.687 846 1
H	4.458 520 6	-2.470 161 8	0.162 253 3
H	5.963 159 2	-1.147 833 5	1.304 242 4
H	7.871 681 5	-0.873 473 7	2.859 289 8
H	8.597 820 0	-5.106 720 5	2.710 912 9
H	6.683 844 8	-5.390 731 7	1.188 342 6
H	4.755 217 8	-3.216 849 8	-2.033 559 9
H	6.356 617 0	-2.714 162 9	-1.463 184 9
H	4.621 552 1	-5.494 873 0	0.577 588 2
H	3.601 461 0	-4.332 010 7	1.437 100 0

Table TS14: B97-D3/aug-cc-pVTZ optimized geometry of the ibuprofen.

At.	X	Y	Z
C	-0.667 998 4	-0.568 767 6	-1.090 282 7
C	0.714 766 0	-0.444 247 8	-0.966 621 3
C	1.329 561 4	-0.528 692 9	0.291 565 0
C	0.522 803 2	-0.738 914 2	1.414 358 2
C	-0.862 634 9	-0.866 118 6	1.284 844 0
C	-1.482 740 7	-0.782 924 8	0.032 315 9
H	-1.126 046 3	-0.497 953 8	-2.075 060 5
H	1.325 508 4	-0.271 768 9	-1.849 556 5
H	0.979 478 6	-0.804 370 9	2.400 063 4
H	-1.471 673 5	-1.028 727 3	2.171 629 0
C	-2.983 364 1	-0.871 859 7	-0.114 953 5
H	-3.229 533 2	-1.554 713 2	-0.939 327 8
H	-3.422 565 6	-1.301 564 7	0.795 546 8
C	-3.665 379 1	0.491 425 5	-0.391 702 9
C	-3.433 439 9	1.482 619 5	0.756 287 6
H	-3.868 712 7	1.097 062 1	1.687 903 3
H	-2.365 972 2	1.652 806 4	0.929 136 2
H	-3.905 018 3	2.447 968 5	0.539 542 5
C	2.845 840 7	-0.423 386 6	0.429 582 4
C	3.300 672 2	0.905 021 0	-0.168 350 8
O	3.205 413 0	1.920 728 6	0.741 328 5
H	3.461 436 4	2.733 776 2	0.273 297 5
O	3.673 440 3	1.079 026 8	-1.307 798 7
C	-5.164 781 0	0.289 316 5	-0.649 728 4
H	-5.652 631 9	-0.134 251 1	0.237 976 0
H	-5.657 445 5	1.241 085 3	-0.879 676 8
H	-5.337 858 4	-0.396 361 6	-1.487 754 7
H	-3.208 957 7	0.907 438 3	-1.301 638 4
C	3.573 719 0	-1.599 969 8	-0.242 110 2
H	4.656 806 2	-1.520 267 0	-0.101 181 4
H	3.226 449 4	-2.542 330 4	0.192 194 5
H	3.370 530 0	-1.610 637 8	-1.315 814 0
H	3.086 648 6	-0.398 625 9	1.497 658 0

Table TS15: B97-D3/aug-cc-pVTZ optimized geometry of the norepinephrine.

At.	X	Y	Z
C	-0.549 399 5	-0.206 398 7	-0.514 674 5
C	0.247 569 1	0.906 345 3	-0.228 350 9
C	1.602 319 5	0.755 452 4	0.064 347 4
C	2.191 381 0	-0.520 849 0	0.080 053 2
C	1.399 294 5	-1.632 907 5	-0.209 658 0
C	0.043 900 6	-1.476 008 5	-0.504 755 6
O	2.447 733 9	1.813 321 3	0.350 545 5
H	1.960 018 8	2.642 600 4	0.276 030 0
O	3.521 587 1	-0.695 755 3	0.364 876 6
H	3.913 988 9	0.174 159 5	0.526 565 1
C	-2.040 623 0	-0.048 001 7	-0.815 079 8
O	-2.440 239 2	1.316 145 4	-0.940 020 6
C	-2.901 774 8	-0.732 460 7	0.269 961 5
N	-2.765 267 7	0.034 782 8	1.519 820 6
H	-1.848 161 2	-0.121 636 6	1.929 036 7
H	-3.460 918 2	-0.244 867 5	2.202 326 6
H	-2.658 847 7	1.591 914 1	-0.032 411 0
H	-0.195 738 8	1.899 217 2	-0.257 361 8
H	1.864 884 2	-2.614 328 4	-0.204 480 6
H	-0.552 248 9	-2.355 472 8	-0.736 976 3
H	-2.615 229 9	-1.791 550 0	0.356 490 9
H	-2.256 783 8	-0.528 066 9	-1.779 095 1
H	-3.949 966 9	-0.684 750 7	-0.045 630 6

Table TS16: B97-D3/aug-cc-pVTZ optimized geometry of the phenethylamine

At.	X	Y	Z
C	2.316 478 1	-0.514 000 0	-6.337 050 1
C	2.613 026 0	-1.135 893 5	-7.556 851 7
C	3.073 238 3	-2.455 007 9	-7.596 442 4
C	3.247 187 8	-3.170 483 9	-6.408 945 8
C	2.955 443 4	-2.557 871 6	-5.186 294 9
C	2.491 804 6	-1.240 841 9	-5.150 767 2
C	1.764 893 8	0.904 098 6	-6.313 555 1
C	0.230 644 8	0.889 545 8	-6.200 564 4
N	2.316 194 7	1.653 699 1	-5.166 405 3
H	3.328 515 5	1.712 969 5	-5.239 016 2
H	1.958 018 0	2.604 738 9	-5.169 239 5
H	2.484 739 7	-0.580 950 6	-8.484 828 5
H	3.301 729 8	-2.920 296 2	-8.552 349 3
H	3.610 670 6	-4.194 701 6	-6.435 529 2
H	3.090 857 7	-3.107 285 7	-4.257 504 5
H	2.268 745 6	-0.759 750 4	-4.202 761 0
H	2.024 036 0	1.372 013 6	-7.280 317 3
H	-0.068 911 7	0.406 317 6	-5.264 826 2
H	-0.209 420 9	0.339 817 3	-7.038 574 4
H	-0.167 785 9	1.912 041 6	-6.209 866 9

Table TS17: B97-D3/aug-cc-pVTZ optimized geometry of the proline.

At.	X	Y	Z
C	1.236 608 3	-0.363 083 0	-6.724 062 8
C	2.419 588 0	-1.286 917 6	-6.386 309 1
C	2.733 952 5	-0.941 562 6	-4.893 955 7
N	2.000 642 0	0.298 100 5	-4.607 178 4
C	1.545 148 7	0.879 041 8	-5.884 413 5
H	2.584 028 6	0.945 169 6	-4.086 849 2
C	4.236 571 9	-0.816 867 2	-4.671 360 8
O	4.841 391 1	0.212 865 4	-4.462 519 2
O	4.847 903 3	-2.036 874 5	-4.736 887 4
H	5.797 529 3	-1.879 928 5	-4.599 539 5
H	1.163 701 4	-0.147 270 4	-7.794 111 5
H	0.292 802 8	-0.811 125 6	-6.393 194 4
H	3.278 459 3	-1.036 944 6	-7.019 115 1
H	2.206 008 6	-2.349 626 2	-6.520 478 4
H	2.385 990 4	-1.753 883 1	-4.240 717 6
H	0.669 006 8	1.516 197 4	-5.720 471 1
H	2.323 523 5	1.484 103 4	-6.382 837 2

 Table TS18: Components¹ of the stabilization energy² as well as thermodynamic parameters for the complexes^{3,4,5} under study.

System	$E_{\text{int}}^{\text{DFT}}$	$E_{\text{int}}^{\text{SAPT}}$	E_{Def}^A	E_{Def}^B	E_{Def}	$\Delta ZPVE$	$E_{\text{Def}} + \Delta ZPVE$	E_{Stab}	ΔH^0	ΔG^0
(R)Ibu – (R)Phe	-64.2	-57.2	24.4	1.9	26.3	2.3	28.6	-28.6	-27.6	22.0
(R)Ibu – (S)Phe	-72.5	-65.8	22.4	7.4	29.8	4.0	33.8	-32.0	-31.7	22.0
(R)Ibu – (R)Pro	-71.7	-64.6	22.9	1.0	23.9	2.7	26.6	-38.0	-38.1	12.6
(R)Ibu – (S)Pro	-74.7	-65.8	23.6	3.7	27.4	5.4	32.7	-33.1	-33.5	20.9
(S)Nor – (S)Phe	-71.3	-64.9	25.2	7.6	32.8	6.0	38.8	-26.1	-26.0	27.2
(S)Nor – (R)Phe	-58.4	-51.3	13.2	7.1	20.3	5.9	26.3	-25.0	-25.2	28.7
(R)Nor – (R)Pro	-56.6	-50.9	18.0	4.6	22.6	6.0	28.6	-22.3	-21.8	28.2
(S)Nor – (R)Pro	-60.8	-56.9	18.9	18.7	37.6	6.4	43.9	-13.0	-13.5	39.3
(S)Bac – (S)Phe	-52.5	-50.0	1.3	7.7	9.0	11.8	20.8	-29.2	-33.8	17.5
(S)Bac – (R)Phe	-87.5	-79.2	35.1	3.1	38.2	8.9	47.1	-32.1	-36.5	15.8
(S)Bac – (S)Pro	-74.7	-71.6	22.8	15.4	38.2	10.5	48.7	-22.9	-27.6	24.5
(R)Bac – (S)Pro	-75.7	-67.2	26.6	4.1	30.6	9.9	40.5	-26.7	-31.0	18.9

¹ The E_{Def}^X , $X = A, B$, is the deformation energy of the monomer X , calculated at the same level, as the DFT interaction energy, i.e. as the difference of the electronic energies of the molecule X with geometry as in the dimer and with geometry optimized for the isolated molecule X . E_{Def} is the sum of deformation energies of monomers A and B . $\Delta ZPVE$ is the difference between ZPVE of the dimer and the sum of ZPVE's for monomers obtained at the same DFT level as before. Finally, E_{Stab} is the sum of $E_{\text{int}}^{\text{SAPT}}$ and the deformation and $\Delta ZPVE$ terms.

² Energies are in kJ/mol.

³ $E_{\text{int}}^{\text{DFT}}$ has been computed at the B97-D3/aug-cc-pVTZ level in the dimer basis set.

⁴ $E_{\text{int}}^{\text{SAPT}}$ has been computed in the CBS limit using aug-cc-pVTZ and aug-cc-pVQZ basis sets.

⁵ Thermodynamic corrections to the ΔH^0 and ΔG^0 have been computed at the B97-D3/aug-cc-pVTZ level (298.15K, 0.1MPa).

Table TS19: IQA contributions to the “interaction” energy¹ between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule, $E_{\text{int}}^{\text{IQA}}$, for **Bac – Phe** complexes^{2,3,4} and their differences (see the last column).

Frag1	Frag2	(S) Bac – (S) Phe	(S) Bac – (R) Phe	Diff
Ring	Ring	−26.4	−37.6	−11.2
Ring	*CH	7.2	3.8	−3.4
Ring	NH ₂	−7.6	−31.3	−23.7
Ring	CH ₃	0.3	0.6	0.3
Cl	Ring	1.0	−11.0	−12.0
Cl	*CH	−6.4	−11.2	−4.8
Cl	NH ₂	6.0	8.9	2.9
Cl	CH ₃	−0.2	−1.6	−1.4
*CH	Ring	−15.5	0.0	15.5
*CH	*CH	6.2	4.6	−1.6
*CH	NH ₂	−18.2	−5.1	13.1
*CH	CH ₃	0.2	0.6	0.4
CH ₂ –x	Ring	−2.5	0.4	2.9
CH ₂ –x	*CH	19.4	12.3	−7.1
CH ₂ –x	NH ₂	−30.7	−13.1	17.6
CH ₂ –x	CH ₃	0.7	1.5	0.8
NH ₂	Ring	1.4	−0.5	−1.9
NH ₂	CH	−16.9	−11.6	5.3
NH ₂	NH ₂	18.1	12.3	−5.8
NH ₂	CH ₃	−0.6	−1.4	−0.8
CH ₂ –y	Ring	−10.1	0.2	10.3
CH ₂ –y	*CH	4.2	2.6	−1.6
CH ₂ –y	NH ₂	−4.4	−8.8	−4.4
CH ₂ –y	CH ₃	0.1	0.3	0.2
COOH	Ring	−16.1	−19.9	−3.8
COOH	*CH	−14.0	8.5	22.5
COOH	NH ₂	−15.7	−107.8	−92.1
COOH	CH ₃	−0.4	−4.4	−3.9
All	All	−121.0	−208.7	−87.7

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

³ CH₂–x – group comprising carbon connected to the NH₂ group

⁴ CH₂–y – group comprising carbon connected to the COOH group

Table TS20: IQA contributions to the “interaction” energy¹ between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule, $E_{\text{int}}^{\text{IQA}}$, for **Bac – Pro** complexes² and their differences (see the last column).

Frag1	Frag2	(S) Bac – (S) Pro	(R) Bac – (S) Pro	Diff
Ring	*Ring	–40.1	–46.9	–6.8
Ring	COOH	–15.8	–6.8	9.0
Cl	*Ring	–27.0	–13.9	13.1
Cl	COOH	0.3	3.0	2.7
*CH	*Ring	2.4	0.8	–1.6
*CH	COOH	–3.9	–1.0	2.9
CH ₂ –x	*Ring	6.1	3.1	–3.0
CH ₂ –x	COOH	–10.0	–3.0	7.0
NH ₂	*Ring	–5.6	–3.0	2.6
NH ₂	COOH	9.9	3.0	–7.0
CH ₂ –y	*Ring	1.9	–13.0	–15.0
CH ₂ –y	COOH	–14.2	–0.8	13.4
COOH	*Ring	–16.8	–92.0	–75.1
COOH	COOH	–62.7	–2.1	60.6
All	All	–175.5	–172.7	2.8

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

Table TS21: IQA contributions to the “interaction” energy¹ between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule, $E_{\text{int}}^{\text{IQA}}$, for **Ibu – Phe** complexes^{2,3,4,5} and their differences (see the last column).

Frag1	Frag2	(R) Ibu – (R) Phe	(R) Ibu – (S) Phe	Diff
Ring	Ring	–37.0	–31.3	5.7
Ring	*CH	–4.9	–24.5	–19.6
Ring	NH ₂	4.2	4.6	0.4
Ring	CH ₃	–12.7	–1.1	11.6
CH ₂	Ring	–5.2	–11.6	–6.4
CH ₂	*CH	1.1	1.2	0.1
CH ₂	NH ₂	–1.2	–1.3	–0.1
CH ₂	CH ₃	0.1	0.1	0.0
CH–x	Ring	–0.4	–0.4	0.0
CH–x	*CH	1.9	2.4	0.5
CH–x	NH ₂	–1.8	–2.0	–0.3
CH–x	CH ₃	0.2	0.1	0.0
CH ₃ –x	Ring	0.0	–0.2	–0.2
CH ₃ –x	*CH	–0.1	–0.2	0.0
CH ₃ –x	NH ₂	0.1	0.1	0.0
CH ₃ –x	CH ₃	0.0	0.0	0.0
CH ₃ –y	Ring	0.0	–0.1	–0.1
CH ₃ –y	*CH	–0.1	–0.1	0.0
CH ₃ –y	NH ₂	0.1	0.1	0.0
CH ₃ –y	CH ₃	0.0	0.0	0.0
*CH	Ring	–0.3	–0.3	0.0
*CH	*CH	1.1	1.4	0.3
*CH	NH ₂	–3.2	–5.3	–2.1
*CH	CH ₃	–5.2	0.1	5.2
CH ₃ –z	Ring	0.1	0.0	–0.1
CH ₃ –z	*CH	3.7	3.9	0.3
CH ₃ –z	NH ₂	–4.6	–4.9	–0.3
CH ₃ –z	CH ₃	0.2	0.2	0.0
COOH	Ring	–4.6	–9.1	–4.5
COOH	*CH	8.6	10.6	2.0
COOH	NH ₂	–88.9	–99.1	–10.2
COOH	CH ₃	0.5	0.4	0.0
All	All	–148.3	–166.2	–17.8

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

³ CH–x – group which is located between the CH₃–x and CH₃–y

⁴ CH – group comprising asymmetric carbon

⁵ CH₃–z – group connected to the asymmetric carbon

Table TS22: IQA contributions to the “interaction” energy between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule¹ $E_{\text{int}}^{\text{IQA}}$, for **Ibu – Pro** complexes² and their differences (see the last column).

Frag1	Frag2	(R) Ibu – (R) Pro	(R) Ibu – (S) Pro	Diff
Ring	*Ring	–48.0	–61.6	–13.5
Ring	COOH	–1.4	–1.4	–0.1
CH ₂	*Ring	1.7	0.3	–1.4
CH ₂	COOH	–6.1	–0.5	5.6
CH–x	*Ring	1.6	1.2	–0.4
CH–x	COOH	–2.1	–0.7	1.4
CH ₃ –x	*Ring	0.0	0.0	0.0
CH ₃ –x	COOH	–0.2	–0.1	0.2
CH ₃ –y	*Ring	–0.1	0.0	0.0
CH ₃ –y	COOH	0.0	0.0	0.0
*CH	*Ring	0.1	–6.4	–6.4
*CH	COOH	–1.0	–0.6	0.4
CH ₃ –z	*Ring	0.6	0.2	–0.4
CH ₃ –z	COOH	–1.1	–0.8	0.3
COOH	*Ring	–104.8	–93.8	11.0
COOH	COOH	0.4	1.9	1.6
All	All	–160.5	–162.2	–1.8

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

Table TS23: IQA contributions to the “interaction” energy¹ between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule, $E_{\text{int}}^{\text{IQA}}$, for **Nor – Phe** complexes^{2,3,4,5} and their differences (see the last column).

Frag1	Frag2	(S) Nor – (S) Phe	(S) Nor – (R) Phe	Diff
Ring	Ring	−27.2	−23.4	3.8
Ring	*CH	69.9	86.7	16.8
Ring	NH ₂	−94.8	−101.7	−6.9
Ring	CH ₃	4.4	4.9	0.4
OH-x	Ring	−9.8	−0.9	8.8
OH-x	*CH	−33.4	−62.1	−28.7
OH-x	NH ₂	−38.3	51.7	90.0
OH-x	CH ₃	−2.5	−3.6	−1.1
OH-y	Ring	−4.7	−9.4	−4.8
OH-y	*CH	−29.5	−31.5	−2.1
OH-y	NH ₂	31.4	−36.5	−68.0
OH-y	CH ₃	−1.9	−1.8	0.1
*CH	Ring	−0.6	−2.3	−1.7
*CH	*CH	17.8	20.9	3.1
*CH	NH ₂	−17.0	−19.0	−2.1
*CH	CH ₃	0.9	0.9	0.0
CH ₂	Ring	−2.6	−3.4	−0.8
CH ₂	*CH	13.8	12.9	−0.8
CH ₂	NH ₂	−12.5	−11.4	1.1
CH ₂	CH ₃	0.7	0.5	−0.1
NH ₂	Ring	−21.0	−16.5	4.5
NH ₂	*CH	−10.8	−11.8	−1.0
NH ₂	NH ₂	9.5	10.4	0.8
NH ₂	CH ₃	−0.5	−0.5	0.0
OH-z	Ring	0.2	1.7	1.5
OH-z	*CH	−18.0	−18.1	−0.1
OH-z	NH ₂	16.8	16.5	−0.3
OH-z	CH ₃	−0.8	−0.7	0.1
All	All	−160.1	−147.6	12.5

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

³ OH-x – group connected to the ring in the *para* position

⁴ OH-y – group connected to the ring in the *meta* position

⁵ OH-z – group connected to the asymmetric carbon

Table TS24: IQA contributions to the “interaction” energy¹ between the fragment (Frag1) of the first molecule and the fragment (Frag2) of the second molecule, $E_{\text{int}}^{\text{IQA}}$, for **Nor – Pro** complexes² and their differences (see the last column).

Frag1	Frag2	(R)Nor – (R)Pro	(S)Nor – (R)Pro	Diff
Ring	*Ring	12.3	2.7	-9.6
Ring	COOH	-56.8	-60.8	-4.0
OH-x	*Ring	-44.2	-35.9	8.2
OH-x	COOH	-21.9	0.8	22.7
OH-y	*Ring	-31.1	-21.9	9.3
OH-y	COOH	13.7	-23.0	-36.7
*CH	*Ring	13.0	12.1	-0.9
*CH	COOH	-9.8	-9.3	0.5
CH ₂	*Ring	4.2	5.8	1.6
CH ₂	COOH	-5.9	-5.0	0.9
NH ₂	*Ring	-20.2	-17.2	3.1
NH ₂	COOH	5.5	4.5	-1.0
OH-z	*Ring	-12.6	-10.2	2.3
OH-z	COOH	9.5	8.2	-1.3
All	All	-144.3	-149.2	-5.0

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

Table TS25: F-SAPT/jun-cc-pVDZ interaction energy partitioning for the **Bac – Phe** complexes.

Frag1	Frag2	(S)Bac – (S)Phe						(S)Bac – (R)Phe						Diff	
		Elst	Exch	IndAB	IndBA	Disp	Total	Elst	Exch	IndAB	IndBA	Disp	Total	Total	Total
Ring	Ring	-5.9	14.3	-1.3	-1.0	-16.4	-10.3	-7.7	25.7	-1.9	0.2	-23.9	-7.6	2.7	
Ring	*CH	0.5	0.0	0.1	0.0	-0.4	0.1	0.7	0.4	0.7	0.5	-2.3	0.0	-0.2	
Ring	NH ₂	-8.6	0.1	-0.4	-0.3	-0.8	-10.0	-10.9	16.0	-2.1	-1.4	-11.5	-9.9	0.1	
Ring	CH ₃	2.1	0.0	0.3	0.0	-0.2	2.2	-6.1	0.2	-0.1	0.3	-0.8	-6.5	-8.7	
C1	Ring	-2.4	0.0	0.0	-0.3	-0.3	-3.1	-6.3	7.3	-0.3	-1.0	-5.7	-6.0	-2.9	
C1	*CH	0.3	0.0	0.0	0.0	0.0	0.2	1.0	0.0	0.1	-0.3	-0.3	0.5	0.2	
C1	NH ₂	-1.2	0.0	0.0	-0.1	0.0	-1.3	0.9	0.0	0.0	0.4	-0.6	0.8	2.0	
C1	CH ₃	0.7	0.0	0.0	0.0	0.0	0.7	1.8	0.0	0.0	-0.1	-0.1	1.6	0.9	
*CH	Ring	2.9	12.9	-0.4	0.8	-7.2	9.0	0.5	0.0	0.0	-0.1	-0.4	0.1	-9.0	
*CH	*CH	-1.2	0.1	-0.1	0.1	-0.6	-1.7	-0.3	0.0	0.0	-0.1	-0.1	-0.4	1.3	
*CH	NH ₂	9.1	7.8	-1.3	0.4	-2.9	13.0	0.0	0.1	-0.1	0.0	-0.6	-0.6	-13.7	
*CH	CH ₃	-3.3	0.0	0.4	0.0	-0.2	-3.1	0.4	0.0	0.0	-0.1	-0.1	0.3	3.3	
CH ₂ -x	Ring	-4.6	0.0	0.0	-0.1	-1.2	-5.9	-0.2	0.0	0.0	0.0	-0.1	-0.3	5.5	
CH ₂ -x	*CH	1.0	0.1	0.0	0.0	-0.3	0.9	0.9	0.0	0.0	-0.1	0.0	0.8	-0.1	
CH ₂ -x	NH ₂	5.4	4.1	-1.0	0.2	-2.1	6.6	-3.1	0.0	0.0	-0.5	-0.2	-3.8	-10.4	
CH ₂ -x	CH ₃	-0.5	0.0	0.2	0.0	-0.1	-0.4	0.1	0.0	0.0	0.0	0.0	0.1	0.5	
NH ₂	Ring	-1.4	0.1	0.0	-0.7	-0.4	-2.4	-0.8	0.0	0.0	-0.1	-0.1	-1.0	1.4	
NH ₂	*CH	-0.6	0.0	0.0	-0.1	-0.1	-0.8	-0.3	0.0	0.0	0.0	0.0	-0.3	0.5	
NH ₂	NH ₂	-8.9	0.0	-0.2	-0.6	-0.3	-10.1	0.8	0.0	0.0	0.2	-0.1	0.9	11.0	
NH ₂	CH ₃	2.0	0.0	0.0	-0.1	0.0	2.0	0.6	0.0	0.0	0.0	0.0	0.6	-1.3	
CH ₂ -y	Ring	-1.2	5.7	-0.2	-0.6	-5.0	-1.3	-1.7	0.1	0.1	0.3	-0.6	-1.9	-0.6	
CH ₂ -y	*CH	-0.2	0.0	0.0	0.0	-0.2	-0.4	-1.1	0.2	0.1	0.3	-0.3	-0.8	-0.4	
CH ₂ -y	NH ₂	-4.5	0.2	-0.3	0.0	-0.9	-5.5	7.3	3.6	-0.6	1.2	-2.2	9.3	14.8	
CH ₂ -y	CH ₃	1.1	0.0	0.1	0.0	-0.1	1.1	-0.4	0.0	0.0	0.1	-0.2	-0.4	-1.4	
COOH	Ring	-16.2	7.9	-0.8	-3.2	-4.9	-17.1	-0.1	14.0	0.4	-2.5	-6.6	5.2	22.3	
COOH	*CH	-1.6	0.3	-0.1	-0.4	-0.7	-2.5	8.1	0.8	1.0	-2.3	-1.7	5.9	8.4	
COOH	NH ₂	-24.3	14.5	-2.1	-2.7	-5.4	-19.9	-123.5	100.0	-10.2	-36.1	-15.0	-84.8	-64.9	
COOH	CH ₃	3.6	0.1	0.1	-0.4	-0.5	2.9	6.2	2.0	0.7	-1.2	-2.2	5.5	2.6	
Ring	All	-11.8	14.4	-1.3	-1.4	-17.9	-17.9	-24.1	42.3	-3.4	-0.4	-38.5	-24.0	-6.1	
Cl	All	-2.6	0.0	0.0	-0.4	-0.4	-3.4	-2.5	7.3	-0.2	-1.0	-6.7	-3.2	0.3	
*CH	All	7.4	20.8	-1.3	1.3	-10.9	17.3	0.6	0.1	-0.1	-0.3	-1.1	-0.7	-18.0	
CH ₂ -x	All	1.3	4.3	-0.8	0.1	-3.7	1.2	-2.3	0.0	0.0	-0.6	-0.4	-3.3	-4.5	
NH ₂	All	-9.0	0.1	-0.2	-1.5	-0.8	-11.3	0.4	0.0	0.0	0.1	-0.2	0.3	11.6	
CH ₂ -y	All	-4.8	6.0	-0.5	-0.6	-6.3	-6.2	4.1	3.9	-0.4	2.0	-3.3	6.2	12.4	
COOH	All	-38.5	22.8	-2.9	-6.6	-11.5	-36.7	-109.3	116.8	-8.2	-42.0	-25.4	-68.3	-31.6	
All	Ring	-28.8	40.9	-2.7	-5.1	-35.4	-31.1	-16.3	47.1	-1.7	-3.2	-37.4	-11.5	19.5	
All	*CH	-1.9	0.6	0.0	-0.4	-2.4	-4.1	9.1	1.4	1.8	-1.9	-4.8	5.5	9.6	
All	NH ₂	-33.0	26.7	-5.3	-3.1	-12.5	-27.1	-128.6	119.7	-13.1	-36.2	-30.1	-88.3	-61.1	
All	CH ₃	5.6	0.1	1.1	-0.4	-1.2	5.3	2.7	2.2	0.6	-0.9	-3.3	1.2	-4.1	
All	All	-58.0	68.3	-6.9	-9.0	-51.4	-57.0	-133.2	170.4	-12.4	-42.3	-75.6	-93.0	-36.0	

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

Table TS26: F-SAPT/jun-cc-pVDZ interaction energy partitioning for the **Bac – Pro** complexes.

Frag1	Frag2	(S)Bac – (S)Pro						(R)Bac – (S)Pro						Diff	
		Elst	Exch	IndAB	IndBA	Disp	Total	Elst	Exch	IndAB	IndBA	Disp	Total	Total	Total
Ring	*Ring	-14.5	27.0	-1.4	0.2	-25.0	-13.9	-7.2	31.3	-0.7	-1.0	-27.1	-4.8	9.0	
Ring	COOH	-12.5	8.2	-4.8	-0.5	-7.8	-17.4	-14.5	1.5	-1.2	0.3	-3.7	-17.7	-0.3	
Cl	*Ring	0.0	7.9	-0.6	-0.5	-7.2	-0.5	-4.4	2.8	-0.1	-0.4	-3.4	-5.4	-4.9	
Cl	COOH	-2.6	0.0	-0.5	-0.2	-0.4	-3.7	5.4	0.0	-0.1	-0.1	-0.5	4.6	8.3	
*CH	*Ring	0.8	0.0	0.0	-0.2	-0.6	0.2	-2.6	0.2	-0.1	-0.5	-1.2	-4.2	-4.3	
*CH	COOH	-2.9	0.2	-0.3	-0.4	-0.5	-3.9	1.7	0.0	0.0	-0.1	-0.1	1.5	5.5	
CH ₂ -x	*Ring	0.4	0.0	0.0	0.0	-0.2	0.2	-2.1	0.0	0.0	-0.5	-0.6	-3.2	-3.4	
CH ₂ -x	COOH	-0.4	0.0	-0.1	-0.3	-0.2	-1.0	0.4	0.0	0.0	0.0	-0.1	0.3	1.4	
NH ₂	*Ring	0.3	0.0	0.0	0.0	-0.1	0.3	0.6	0.0	0.0	0.3	-0.2	0.7	0.5	
NH ₂	COOH	1.3	0.0	0.0	0.2	-0.1	1.4	1.2	0.0	0.0	0.0	0.0	1.1	-0.2	
CH ₂ -y	*Ring	-1.9	0.1	0.2	0.5	-0.9	-2.0	7.3	8.0	-0.8	2.5	-5.4	11.4	13.4	
CH ₂ -y	COOH	12.8	7.4	-1.7	1.8	-3.2	17.1	-3.4	0.0	0.1	0.2	-0.3	-3.3	-20.4	
COOH	*Ring	4.0	10.1	0.4	-2.2	-6.0	6.3	-101.3	83.4	-7.0	-32.4	-18.0	-75.3	-81.6	
COOH	COOH	-75.8	40.4	-4.4	-15.6	-9.1	-64.5	18.3	1.6	1.2	-1.1	-2.8	17.3	81.8	
Ring	All	-27.0	35.1	-6.3	-0.3	-32.8	-31.3	-21.8	32.8	-2.0	-0.8	-30.8	-22.5	8.8	
Cl	All	-2.7	7.9	-1.1	-0.8	-7.6	-4.1	1.0	2.9	-0.3	-0.5	-3.9	-0.8	3.4	
*CH	All	-2.1	0.2	-0.3	-0.5	-1.1	-3.8	-0.8	0.2	-0.1	-0.5	-1.3	-2.6	1.1	
CH ₂ -x	All	0.0	0.0	-0.1	-0.3	-0.4	-0.8	-1.7	0.0	0.0	-0.5	-0.7	-2.9	-2.0	
NH ₂	All	1.6	0.0	0.0	0.2	-0.2	1.6	1.8	0.0	0.0	0.3	-0.2	1.9	0.2	
CH ₂ -y	All	11.0	7.5	-1.5	2.3	-4.2	15.1	3.9	8.0	-0.7	2.7	-5.7	8.1	-7.0	
COOH	All	-71.8	50.5	-4.0	-17.8	-15.1	-58.2	-83.0	85.0	-5.8	-33.5	-20.7	-58.0	0.2	
All	*Ring	-10.9	45.1	-1.4	-2.2	-40.0	-9.4	-109.6	125.7	-8.8	-32.0	-55.9	-80.7	-71.3	
All	COOH	-80.1	56.3	-11.9	-15.1	-21.3	-72.1	9.1	3.2	-0.1	-0.8	-7.5	3.9	76.0	
All	All	-91.0	101.3	-13.3	-17.2	-61.3	-81.5	-100.5	128.9	-8.9	-32.8	-63.4	-76.8	4.7	

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

 Table TS27: F-SAPT/jun-cc-pVDZ interaction energy partitioning for the **Ibu – Phe** complexes.

Frag1	Frag2	(R)Ibu – (R)Phe						(R)Ibu – (S)Phe						Diff	
		Elst	Exch	IndAB	IndBA	Disp	Total	Elst	Exch	IndAB	IndBA	Disp	Total	Total	Total
Ring	Ring	-3.9	28.0	-1.3	-1.5	-25.5	-4.3	-14.0	21.7	-1.8	0.1	-27.1	-21.1	-16.8	
Ring	*CH	-2.1	0.2	0.2	0.0	-1.1	-2.8	-4.2	14.3	0.0	0.4	-7.3	3.1	6.0	
Ring	NH ₂	-0.3	0.2	0.0	0.1	-1.4	-1.4	13.5	1.8	-1.2	2.1	-3.5	12.6	14.0	
Ring	CH ₃	-4.0	8.0	-0.3	-0.1	-5.1	-1.5	-2.9	0.1	0.5	0.1	-1.4	-3.5	-2.0	
CH ₂	Ring	1.3	2.0	-0.1	-0.3	-3.3	-0.4	-1.1	6.8	-0.2	0.1	-5.4	-0.1	0.4	
CH ₂	*CH	0.0	0.0	0.0	-0.1	0.0	-0.1	-1.3	0.0	0.0	-0.1	-0.2	-1.5	-1.4	
CH ₂	NH ₂	-1.3	0.0	0.0	-0.2	0.0	-1.5	-1.7	0.0	0.0	-0.3	-0.1	-2.1	-0.6	
CH ₂	CH ₃	-0.2	0.0	0.0	-0.1	-0.1	-0.3	1.0	0.0	0.0	0.0	-0.1	0.9	1.2	
CH-x	Ring	-1.8	0.0	0.0	0.1	-0.6	-2.2	-0.1	0.1	0.0	0.0	-0.9	-0.9	1.3	
CH-x	*CH	0.2	0.0	0.0	0.0	0.0	0.2	-0.4	0.0	0.0	0.1	0.0	-0.3	-0.5	
CH-x	NH ₂	0.6	0.0	0.0	0.0	0.0	0.6	1.0	0.0	0.0	0.1	0.0	1.1	0.5	
CH-x	CH ₃	-0.1	0.0	0.0	0.0	0.0	-0.1	-0.1	0.0	0.0	0.0	0.0	-0.1	0.0	
CH ₃ -y	Ring	-0.2	0.0	0.0	0.2	-0.3	-0.3	-1.6	0.1	0.0	0.1	-0.3	-1.8	-1.5	
CH ₃ -x	*CH	0.1	0.0	0.0	0.0	0.1	0.1	0.3	0.0	0.0	0.0	0.0	0.3	0.2	
CH ₃ -x	NH ₂	-0.1	0.0	0.0	0.0	0.0	-0.1	0.1	0.0	0.0	0.0	0.0	0.1	0.2	
CH ₃ -x	CH ₃	0.3	0.0	0.0	0.0	0.0	0.3	0.2	0.0	0.0	0.0	0.0	0.2	-0.1	
CH ₃ -y	Ring	-0.6	0.0	0.0	0.1	-0.3	-0.8	-1.2	0.0	0.0	0.1	-0.4	-1.6	-0.8	
CH ₃ -y	*CH	0.4	0.0	0.0	0.0	0.0	0.4	-0.6	0.0	0.0	0.0	0.0	-0.6	-1.0	
CH ₃ -y	NH ₂	-0.5	0.0	0.0	-0.1	0.0	-0.5	-0.2	0.0	0.0	-0.1	0.0	-0.3	0.3	
CH ₃ -y	CH ₃	-0.1	0.0	0.0	0.0	0.0	-0.1	0.6	0.0	0.0	0.0	0.0	0.6	0.7	
*CH	Ring	-4.9	0.1	0.1	0.7	-1.1	-5.0	7.2	0.0	0.0	-0.2	-0.8	6.3	11.4	
*CH	*CH	-1.8	0.1	0.0	0.3	-0.3	-1.7	12.7	0.0	0.0	-0.5	-0.5	11.7	13.4	
*CH	NH ₂	11.2	1.4	-0.4	1.6	-1.1	12.8	-5.2	2.2	-0.5	0.2	-1.5	-4.8	-17.6	
*CH	CH ₃	0.5	2.3	0.0	0.4	-1.6	1.6	-8.1	0.0	0.1	0.1	-0.1	-8.1	-9.6	
CH ₃ -z	Ring	1.4	0.0	0.0	-0.1	-0.4	0.9	0.9	0.0	0.0	-0.1	-0.3	0.6	-0.4	
CH ₃ -z	*CH	0.9	0.0	0.0	-0.1	-0.1	0.7	-1.3	0.0	0.0	-0.2	-0.1	-1.5	-2.3	
CH ₃ -z	NH ₂	-6.0	0.0	-0.1	-0.8	-0.2	-7.0	-6.7	0.1	-0.1	-1.0	-0.3	-8.1	-1.1	
CH ₃ -z	CH ₃	0.4	0.0	0.0	-0.2	-0.3	0.0	2.0	0.0	0.0	-0.1	0.0	1.9	1.9	
COOH	Ring	8.8	2.5	0.9	-1.4	-3.3	7.6	7.5	5.7	1.0	-1.6	-4.8	7.8	0.1	
COOH	*CH	10.6	1.2	0.9	-1.6	-1.2	9.9	-0.5	0.6	0.4	-1.6	-1.4	-2.5	-12.4	
COOH	NH ₂	-98.0	70.4	-7.2	-23.9	-11.8	-70.5	-109.3	83.5	-8.2	-29.9	-13.2	-77.1	-6.6	
COOH	CH ₃	1.0	0.1	0.4	-0.9	-0.8	-0.2	10.4	0.3	1.0	-0.4	-0.4	10.9	11.2	
Ring	All	-10.3	36.4	-1.4	-1.5	-33.2	-10.0	-7.6	37.9	-2.5	2.6	-39.3	-8.9	1.1	
CH ₂	All	-0.3	2.0	-0.1	-0.6	-3.4	-2.4	-3.1	6.9	-0.2	-0.5	-5.7	-2.7	-0.3	
CH-x	All	-1.1	0.0	0.0	0.2	-0.7	-1.5	0.4	0.1	0.0	0.2	-1.0	-0.3	1.2	
CH ₃ -x	All	0.1	0.0	0.0	0.2	-0.3	0.0	-1.0	0.1	0.0	0.1	-0.4	-1.2	-1.2	
CH ₃ -y	All	-0.8	0.0	0.0	0.0	-0.3	-1.0	-1.4	0.0	0.0	0.0	-0.5	-1.8	-0.8	
*CH	All	5.1	3.8	-0.3	3.1	-4.1	7.6	6.6	2.3	-0.4	-0.4	-3.0	5.2	-2.4	
CH ₃ -z	All	-3.2	0.1	0.0	-1.2	-1.0	-5.4	-5.0	0.1	-0.1	-1.3	-0.8	-7.2	-1.8	
COOH	All	-77.7	74.2	-4.9	-27.7	-17.1	-53.2	-91.8	90.1	-5.9	-33.5	-19.8	-60.9	-7.7	
All	Ring	0.2	32.6	-0.4	-2.1	-34.8	-4.4	-2.4	34.4	-1.0	-1.7	-40.1	-10.9	-6.4	
All	*CH	8.3	1.4	1.1	-1.4	-2.8	6.6	4.8	15.0	0.3	-1.8	-9.6	8.7	2.1	
All	NH ₂	-94.4	72.0	-7.6	-23.1	-14.6	-67.7	-108.5	87.6	-10.1	-28.9	-18.6	-78.6	-10.9	
All	CH ₃	-2.2	10.5	0.0	-0.8	-7.9	-0.5	3.1	0.5	1.6	-0.3	-2.0	2.8	3.3	
All	All	-88.2	116.5	-6.8	-27.4	-60.2	-66.0	-102.9	137.4	-9.2	-32.8	-70.4	-77.9	-11.8	

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

Table TS28: F-SAPT/jun-cc-pVDZ interaction energy partitioning for the **Ibu – Pro** complexes.

Frag1	Frag2	(R)Ibu – (R)Pro						(R)Ibu – (S)Pro						Diff	
		Elst	Exch	IndAB	IndBA	Disp	Total	Elst	Exch	IndAB	IndBA	Disp	Total	Total	
Ring	*Ring	-8.9	36.2	-3.2	2.6	-23.8	2.8	-3.5	39.6	-1.8	3.4	-30.5	7.2	4.4	
Ring	COOH	-7.2	2.9	0.3	0.2	-4.5	-8.4	-3.4	0.9	-0.8	0.2	-3.4	-6.5	1.8	
CH ₂	*Ring	-2.9	0.0	0.2	-0.6	-0.8	-4.1	-1.4	0.0	0.0	-0.6	-1.0	-3.0	1.1	
CH ₂	COOH	10.1	1.6	-0.7	0.0	-1.7	9.3	1.8	0.0	-0.1	0.0	-0.3	1.4	-7.9	
CH-x	*Ring	1.5	0.0	0.0	0.2	-0.2	1.6	0.2	0.0	0.0	0.3	-0.3	0.3	-1.3	
CH-x	COOH	-3.0	0.1	-0.1	0.0	-0.3	-3.3	0.6	0.0	0.0	0.0	-0.1	0.5	3.8	
CH ₃ -x	*Ring	1.0	0.0	0.0	0.1	-0.1	0.9	0.0	0.0	0.0	0.1	-0.2	0.0	-0.9	
CH ₃ -x	COOH	-2.4	0.0	0.0	0.0	-0.1	-2.5	-0.2	0.0	0.0	0.0	0.0	-0.2	2.3	
CH ₃ -y	*Ring	0.1	0.0	0.0	-0.1	-0.1	-0.1	-0.7	0.0	0.0	0.0	-0.1	-0.8	-0.7	
CH ₃ -y	COOH	-0.7	0.0	-0.1	0.0	-0.1	-0.9	0.8	0.0	0.0	0.0	0.0	0.7	1.7	
*CH	*Ring	5.7	0.4	-0.2	1.4	-1.6	5.7	5.9	3.1	-0.4	-0.9	-3.6	4.0	-1.7	
*CH	COOH	-1.9	0.0	0.1	0.0	-0.1	-1.9	-16.3	0.0	0.0	0.2	-0.2	-16.2	-14.3	
CH ₃ -z	*Ring	-5.6	0.0	-0.1	-1.4	-0.5	-7.4	-6.3	0.1	-0.1	-1.6	-0.8	-8.7	-1.3	
CH ₃ -z	COOH	1.8	0.0	0.0	0.0	0.0	1.7	2.6	0.0	0.0	-0.1	-0.1	2.4	0.7	
COOH	*Ring	-109.1	99.7	-8.0	-40.8	-20.3	-78.5	-106.5	91.0	-7.9	-36.9	-17.8	-78.1	0.4	
COOH	COOH	8.1	0.1	1.4	-0.3	-0.7	8.6	20.5	0.2	1.5	-0.6	-1.2	20.4	11.8	
Ring	All	-16.1	39.1	-2.9	2.8	-28.3	-5.6	-6.9	40.5	-2.6	3.6	-34.0	0.7	6.2	
CH ₂	All	7.2	1.6	-0.5	-0.6	-2.5	5.3	0.4	0.0	0.0	-0.6	-1.4	-1.6	-6.8	
CH-x	All	-1.4	0.1	-0.1	0.2	-0.5	-1.7	0.8	0.0	0.0	0.3	-0.3	0.8	2.5	
CH ₃ -x	All	-1.5	0.0	0.0	0.1	-0.2	-1.6	-0.2	0.0	0.0	0.2	-0.2	-0.2	1.4	
CH ₃ -y	All	-0.6	0.0	0.0	-0.1	-0.2	-1.0	0.1	0.0	0.0	-0.2	0.0	0.0	1.0	
*CH	All	3.8	0.4	-0.1	1.4	-1.7	3.8	-10.4	3.1	-0.4	-0.7	-3.8	-12.2	-16.0	
CH ₃ -z	All	-3.8	0.0	0.0	-1.4	-0.5	-5.7	-3.7	0.1	-0.1	-1.7	-0.9	-6.2	-0.6	
COOH	All	-101.0	99.8	-6.6	-41.1	-20.9	-69.9	-85.9	91.2	-6.4	-37.5	-19.0	-57.7	12.2	
All	*Ring	-118.2	136.3	-11.2	-38.5	-47.4	-79.0	-112.2	133.8	-10.2	-36.1	-54.3	-79.0	0.0	
All	COOH	4.9	4.8	0.9	-0.2	-7.6	2.7	6.4	1.2	0.7	-0.4	-5.4	2.5	-0.2	
All	All	-113.3	141.1	-10.3	-38.8	-55.0	-76.3	-105.8	135.0	-9.5	-36.5	-59.7	-76.5	-0.2	

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

Table TS29: F-SAPT/jun-cc-pVDZ interaction energy partitioning for the **Nor – Phe** complexes.

Frag1	Frag2	(S)Nor – (S)Phe						(S)Nor – (R)Phe						Diff	
		Elst	Exch	IndAB	IndBA	Disp	Total	Elst	Exch	IndAB	IndBA	Disp	Total	Total	
Ring	Ring	13.5	22.3	-1.3	0.0	-25.0	9.6	17.4	15.8	-1.0	-0.4	-19.5	12.3	2.8	
Ring	*CH	-6.7	0.2	0.1	-0.2	-1.3	-7.8	-5.7	0.1	-0.1	0.0	-1.4	-7.1	0.7	
Ring	NH ₂	-13.7	4.9	-2.0	-1.7	-4.5	-17.0	-16.2	0.6	-1.1	-1.6	-2.5	-20.8	-3.8	
Ring	CH ₃	1.1	0.0	0.7	0.0	-0.4	1.3	1.5	0.0	0.5	0.0	-0.4	1.6	0.2	
OH-x	Ring	-6.2	6.4	0.9	-1.7	-5.2	-5.8	-14.2	1.5	-0.1	0.2	-3.2	-15.8	-10.0	
OH-x	*CH	8.6	1.6	0.2	-1.2	-1.9	7.3	12.5	6.7	-0.1	-0.4	-2.3	16.5	9.2	
OH-x	NH ₂	-82.2	80.0	-7.9	-23.1	-11.7	-44.8	6.8	2.7	-0.5	1.0	-1.7	8.2	53.0	
OH-x	CH ₃	3.5	0.1	0.9	-0.2	-0.4	4.0	-7.9	0.2	0.1	0.1	-0.4	-7.9	-11.8	
OH-y	Ring	-6.1	0.5	-0.1	-0.4	-2.1	-8.2	-3.6	6.5	1.1	-0.5	-4.6	-1.2	7.0	
OH-y	*CH	2.5	0.0	0.0	-0.1	-0.1	2.3	0.7	0.5	0.2	-0.5	-1.2	-0.3	-2.6	
OH-y	NH ₂	-0.4	0.0	-0.1	0.0	-0.1	-0.6	-76.0	74.6	-7.4	-21.5	-10.9	-41.1	-40.5	
OH-y	CH ₃	-0.6	0.0	0.0	0.0	0.0	-0.6	6.5	0.2	0.9	-0.2	-0.3	7.2	7.8	
*CH	Ring	-1.8	0.0	0.0	0.1	-0.7	-2.5	-1.5	0.0	0.0	-0.1	-0.6	-2.2	0.3	
*CH	*CH	0.4	0.0	0.0	0.0	0.0	0.4	0.1	0.0	0.0	0.0	0.0	0.1	-0.4	
*CH	NH ₂	-0.5	0.0	0.0	-0.1	0.0	-0.6	-0.2	0.0	0.0	-0.1	0.0	-0.3	0.3	
*CH	CH ₃	0.4	0.0	0.0	0.0	0.0	0.4	0.5	0.0	0.0	0.0	0.5	0.0	0.0	
CH ₂	Ring	-1.3	1.0	-0.1	-0.3	-2.4	-3.0	-1.6	0.5	0.0	-0.1	-1.9	-3.1	-0.1	
CH ₂	*CH	0.2	0.0	0.0	0.0	0.0	0.2	-0.3	0.0	0.0	0.0	0.0	-0.3	-0.5	
CH ₂	NH ₂	-0.4	0.0	0.0	0.0	0.0	-0.5	0.2	0.0	0.0	0.0	0.0	0.2	0.6	
CH ₂	CH ₃	0.1	0.0	0.0	0.0	0.0	0.1	0.1	0.0	0.0	0.0	0.0	0.1	0.0	
NH ₂	Ring	-8.6	10.5	-0.3	-1.0	-6.5	-5.9	-4.8	9.3	-0.2	-0.6	-5.9	-2.2	3.6	
NH ₂	*CH	-0.1	0.0	0.0	0.0	0.0	-0.2	0.4	0.0	0.0	0.0	0.0	0.4	0.6	
NH ₂	NH ₂	-0.9	0.0	0.0	-0.1	0.0	-1.2	-1.0	0.0	0.0	-0.1	0.0	-1.1	0.1	
NH ₂	CH ₃	0.5	0.0	0.0	0.0	0.0	0.5	0.4	0.0	0.0	0.0	0.0	0.4	-0.1	
OH-z	Ring	-1.5	0.0	0.0	0.2	-0.3	-1.6	-1.6	0.0	0.0	-0.1	-0.3	-1.9	-0.4	
OH-z	*CH	-0.1	0.0	0.0	0.0	0.0	-0.1	0.5	0.0	0.0	0.0	0.0	0.5	0.6	
OH-z	NH ₂	-0.9	0.0	0.0	-0.2	0.0	-1.1	-0.8	0.0	0.0	-0.1	0.0	-0.9	0.2	
OH-z	CH ₃	0.6	0.0	0.0	0.0	0.0	0.6	0.5	0.0	0.0	0.0	0.0	0.5	-0.1	
Ring	All	-5.9	27.5	-2.5	-1.9	-31.2	-14.0	-3.1	16.5	-1.7	-2.0	-23.8	-14.1	-0.1	
OH-x	All	-76.2	88.2	-5.9	-26.2	-19.2	-39.3	-2.8	11.1	-0.6	0.8	-7.6	1.0	40.4	
OH-y	All	-4.7	0.5	-0.1	-0.5	-2.3	-7.1	-72.4	81.8	-5.2	-22.6	-17.0	-35.4	-28.3	
*CH	All	-1.4	0.0	0.0	-0.1	-0.8	-2.3	-1.2	0.0	0.0	-0.1	-0.6	-2.0	0.3	
CH ₂	All	-1.4	1.0	-0.1	-0.4	-2.4	-3.2	-1.6	0.5	-0.1	-0.1	-1.9	-3.2	0.1	
NH ₂	All	-9.1	10.5	-0.3	-1.2	-6.6	-6.7	-4.9	9.3	-0.2	-0.7	-6.0	-2.5	4.2	
OH-z	All	-1.8	0.0	0.0	0.0	-0.3	-2.1	-1.3	0.0	0.0	-0.2	-0.3	-1.9	0.2	
All	Ring	-11.9	40.8	-0.8	-3.1	-42.3	-17.3	-10.0	33.7	-0.3	-1.6	-35.9	-14.2	3.2	
All	*CH	4.8	1.9	0.4	-1.5	-3.4	2.1	8.2	7.3	0.0	-0.9	-4.9	9.7	7.6	
All	NH ₂	-99.0	85.0	-10.1	-25.3	-16.4	-65.8	-87.2	77.9	-9.1	-22.4	-15.2	-55.9	9.9	
All	CH ₃	5.7	0.1	1.7	-0.3	-0.9	6.3	1.7	0.4	1.6	-0.1	-1.1	2.4	-4.0	
All	All	-100.5	127.8	-8.9	-30.2	-62.9	-74.7	-87.4	119.3	-7.8	-24.9	-57.2	-58.0	16.7	

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

Table TS30: F-SAPT/jun-cc-pVDZ interaction energy partitioning for the **Nor – Pro** complexes.

Frag1	Frag2	(R)Nor – (R)Pro						(S)Nor – (R)Pro						Diff	
		Elst	Exch	IndAB	IndBA	Disp	Total	Elst	Exch	IndAB	IndBA	Disp	Total	Total	Total
Ring	*Ring	-10.9	27.8	-1.0	-0.4	-24.1	-8.7	-11.9	27.7	-0.9	-1.1	-23.7	-9.9	-1.2	
Ring	COOH	-12.4	3.7	-1.7	-1.0	-4.8	-16.3	-18.6	0.9	-1.9	-1.6	-3.7	-24.8	-8.5	
OH-x	*Ring	-3.4	10.0	-0.7	-0.6	-5.0	0.2	0.1	4.8	0.0	-0.8	-3.8	0.3	0.1	
OH-x	COOH	-40.2	37.1	-3.9	-10.1	-7.6	-24.7	-20.0	10.5	-1.7	-3.4	-4.0	-18.5	6.2	
OH-y	*Ring	1.2	2.9	-0.2	-0.7	-3.3	-0.1	5.0	0.9	0.7	0.0	-2.4	4.2	4.3	
OH-y	COOH	-4.1	0.0	-0.1	-0.7	-0.2	-5.1	-33.1	40.2	-3.9	-10.8	-8.4	-16.1	-11.0	
*CH	*Ring	0.5	0.0	0.0	0.1	-0.7	-0.1	3.5	0.0	0.0	0.8	-0.6	3.8	3.9	
*CH	COOH	0.7	0.0	0.0	0.0	0.0	0.6	-3.9	0.0	0.0	-0.4	0.0	-4.3	-4.9	
CH ₂	*Ring	-0.2	1.4	-0.1	-0.2	-2.1	-1.2	-2.6	0.2	0.0	-0.6	-1.4	-4.4	-3.2	
CH ₂	COOH	-0.7	0.0	0.0	-0.1	-0.1	-0.8	5.1	0.0	0.0	0.4	0.0	5.5	6.3	
NH ₂	*Ring	-1.2	4.8	0.0	0.0	-4.5	-1.0	-0.7	4.1	0.0	-0.3	-3.6	-0.6	0.4	
NH ₂	COOH	1.4	0.0	0.0	0.0	-0.1	1.2	-0.4	0.0	0.0	-0.2	0.0	-0.6	-1.8	
OH-z	*Ring	0.2	0.0	0.0	-0.2	-0.3	-0.2	0.6	0.0	0.0	-0.1	-0.3	0.3	0.6	
OH-z	COOH	0.6	0.0	0.0	0.1	0.0	0.7	0.7	0.0	0.0	-0.1	0.0	0.6	-0.1	
Ring	All	-23.4	31.5	-2.7	-1.4	-28.9	-25.0	-30.5	28.7	-2.7	-2.7	-27.4	-34.6	-9.7	
OH-x	All	-43.6	47.1	-4.6	-10.6	-12.6	-24.4	-19.9	15.3	-1.6	-4.1	-7.8	-18.2	6.3	
OH-y	All	-2.9	2.9	-0.2	-1.4	-3.5	-5.2	-28.1	41.1	-3.3	-10.8	-10.8	-11.9	-6.7	
*CH	All	1.2	0.0	0.0	0.1	-0.8	0.6	-0.3	0.0	0.0	0.4	-0.6	-0.5	-1.1	
CH ₂	All	-0.9	1.4	-0.1	-0.3	-2.2	-2.1	2.5	0.2	0.0	-0.2	-1.4	1.1	3.1	
NH ₂	All	0.1	4.8	-0.1	-0.1	-4.6	0.2	-1.1	4.1	-0.1	-0.5	-3.6	-1.2	-1.4	
OH-z	All	0.8	0.0	0.0	-0.1	-0.3	0.4	1.3	0.0	0.0	-0.2	-0.3	0.9	0.5	
All	*Ring	-13.9	46.9	-2.0	-2.0	-40.1	-11.1	-6.0	37.7	-0.2	-2.0	-35.7	-6.2	4.9	
All	COOH	-54.8	40.9	-5.8	-11.8	-12.8	-44.4	-70.1	51.7	-7.6	-16.0	-16.2	-58.2	-13.8	
All	All	-68.7	87.8	-7.8	-13.8	-52.9	-55.5	-76.1	89.4	-7.8	-18.0	-51.9	-64.5	-9.0	

¹ Energies are in kJ/mol.

² Star (*) denotes the asymmetric carbon.

1 Characteristic features of the monomers' IR spectra

Figure FS1 shows the computed IR spectra for the monomers.

In the spectrum of **Phe** two areas can be distinguished. The region up to ca. 1600 cm^{-1} is composed of peaks corresponding to the rocking movements of CH_3 and NH_2 groups (ca. $255 - 282\text{ cm}^{-1}$), aromatic C-H wagging (700 cm^{-1}), C-H and N-H wagging (826 cm^{-1}), C-C-H bending (1375 cm^{-1}), and NH_2 scissoring (1623 cm^{-1}). Next, in the region between $2875 - 3135\text{ cm}^{-1}$ peaks related to the C-H bonds stretching occur. N-H stretching vibrations are manifested by the small peak at ca. 3420 cm^{-1} .

The spectrum of **Pro** has a different shape. The breathing movements of the pyrrolidine ring give rise to the broad band emerging between $400 - 870\text{ cm}^{-1}$. Next, the high peak at 1072 cm^{-1} is due to the coupled vibration of the ring with the O-H wagging. The sharp peak at 1743 cm^{-1} is related to the C=O stretching. The C-H bonds' stretching vibrations are found between $2980 - 3060\text{ cm}^{-1}$, while the N-H and O-H stretchings can be observed at 3475 and 3653 cm^{-1} , respectively.

The spectrum for ibuprofen comprises five major peaks: (i) the broad, nonsymmetric band at ca. 600 cm^{-1} due to the wagging of OH from the carboxyl group; (ii) the double peak at ca. 1100 cm^{-1} resulting from the aliphatic C-H rocking; (iii) the high peak at 1746 cm^{-1} assigned to the C=O stretching; (iv) the double peak at ca. 3000 cm^{-1} due to the C-H stretching; (v) the 3651 cm^{-1} peak of O-H stretching. The spectrum is in line with the experimental IR ibuprofen spectra available in the literature.^{1,2}

The IR spectrum of the isolated norepinephrine is similar to the experimental and simulated spectra for different conformers described in Ref.³ Here, the whole energy range can be divided into three regions. In the broadest one, up to ca. 1600 cm^{-1} , one can distinguish three highest peaks: (i) at 264 cm^{-1} due to the out-of-plane bending of O-H bonds substituting the phenyl ring; (ii) at 562 cm^{-1} due to the wagging of the OH group connected to the chiral carbon; and (iii) 1236 cm^{-1} due to the C-C-H scissoring. Next, the C-H bonds' stretching motions give rise to the broad band at ca. 3000 cm^{-1} , while the double band between ca. $3600 - 3750\text{ cm}^{-1}$ is due to all the O-H stretching vibrations present in the molecule.

The spectrum computed for baclofen is also characterized by four major peaks: (i) the 966 cm^{-1} peak results from the O-H wagging; (ii) the 1426 cm^{-1} peak results from the C-O-H scissoring; (iii) the 1767 cm^{-1} peak results from the C=O stretching; and (iv) the strongest one at 3159 cm^{-1} due to the O-H stretching (note that on the left shoulder smaller peaks at ca. 3000 cm^{-1} of the C-H stretching emerge). Their positions are comparable to the experimental ones listed in.⁴ The symmetric and antisymmetric stretchings of the NH_2 group give rise to the peak of very low intensity at ca. 3500 cm^{-1} .

2 Characteristic features of the VCD spectra

Table TS31: Assignment of the VCD spectrum of the (R)**Ibu** – (R)**Phe** pair.

peak position [cm ⁻¹]	sign	assignment
589	-	Ibu ring deformation (C–H out of plane bending)
934	+	Ibu O–H wagging coupled to Phe NH ₂ symmetric bend
1004	-	Ibu O–H wagging coupled to Phe *C–H wagging
1185	+	Ibu in plane C–H ring scissoring
1337	-	Ibu *C–H scissoring
1749	-	C=O stretching
3096	-	O–H stretching

Table TS32: Assignment of the VCD spectrum of the (R)**Ibu** – (S)**Phe** pair.

peak position [cm ⁻¹]	sign	assignment
578	-	Ibu ring deformation (C–H out of plane bending)
1073	+	Ibu *C–CH ₃ wagging
1191	+	Phe *C–H wagging coupled to Ibu in plane C–H ring scissoring
1341	+	Ibu *C–H wagging coupled to Phe in plane C–H ring scissoring
1420	+	Ibu O–H coupled to Phe in plane C–H ring scissoring
1751	-	C=O stretching
2979	-	Ibu O–H and CH ₃ stretching coupled to Phe *C–H stretching
2982	+	Ibu O–H and CH ₃ stretching coupled to Phe *C–H stretching

Table TS33: Assignment of the VCD spectrum of the (S)**Nor** – (S)**Phe** pair.

peak position [cm ⁻¹]	sign	assignment
336	+	Nor O–H wagging
344	-	Nor O–H wagging
359	-	Nor O–H wagging
387	-	Nor N–H wagging
393	+	Phe N–H wagging
952	+	Nor in plane C–H ring scissoring coupled to C–C stretching
968	-	Nor O–H wagging coupled to Phe N–N symmetric bend
982	-	Nor O–H wagging coupled to Phe N–N and *C–C ₃ symmetric bend
1589	-	Nor ring deformations due to C–O–H bending
3047	+	Phe CH ₃ stretching
3063	+	Nor O–H stretching (the OH group forming H-bond with Phe) coupled to Nor C–H stretching in ring
3068	+	Nor O–H stretching (the OH group forming H-bond with Phe) coupled to Nor C–H stretching in ring

Table TS34: Assignment of the VCD spectrum of the (S)**Nor** – (R)**Phe** pair.

peak position [cm ⁻¹]	sign	assignment
330	-	both NH ₂ rocking
752	-	Nor ring C–H out of plane bending
848	-	Nor O–H wagging (the OH group forming H-bond with Phe)
1301	-	Nor *C–H wagging
1309	-	Nor *C–H wagging coupled to wagging of *C–CH ₂
3104	+	Phe ring C–H stretching coupled to Nor O–H stretching (the OH group forming H-bond with Phe)
3106	+	Phe ring C–H stretching coupled to Nor O–H stretching (the OH group forming H-bond with Phe)

 Table TS35: Assignment of the VCD spectrum of the (S)**Bac** – (S)**Phe** pair.

peak position [cm ⁻¹]	sign	assignment
428	-	Bac NH ₂ rocking
574	-	Bac skeletal movements
828	+	Bac ring deformation (C–H out of plane bending)
1012	+	Bac O–H wagging
1209	+	Phe C–H ring scissoring coupled to NH ₂ rocking
1288	+	Bac CH ₂ rocking
1465	-	Bac CH ₂ scissoring coupled to C–O–H bending
1626	+	Bac NH ₂ scissoring
3028	+	Bac C–H stretching coupled to O–H stretching

 Table TS36: Assignment of the VCD spectrum of the (S)**Bac** – (R)**Phe** pair.

peak position [cm ⁻¹]	sign	assignment
260	+	Bac NH ₂ rocking
381	-	Bac ring deformation
530	+	Bac ring deformation (C–H out of plane bending)
606	-	Bac CH ₂ rocking
1075	-	Bac O–H wagging
1236	+	Bac *C–H wagging
1301	-	Bac ring C–H rocking coupled to CH ₂ rocking
1424	+	Bac C–O–H bending
1465	-	Phe *C–CH ₃ rocking
2820	+	Bac O–H stretching (the OH group forming H-bond with Phe)
2991	+	Phe C–H stretching

Table TS37: Assignment of the VCD spectrum of the (R)**Ibu** – (R)**Pro** pair.

peak position [cm ⁻¹]	sign	assignment
422	+	skeletal Pro movements
610	-	Pro O–H wagging
817	+	Pro N–H rocking
908	+	Pro ring C ₂ rocking
1045	-	Ibu O–H wagging
1083	+	Ibu O–H wagging coupled to Pro N–H bending
1367	+	Ibu C–O–H bending coupled to Pro CH ₂ rocking
1446	-	Pro N–H wagging
2786	+	Ibu O–H stretching

 Table TS38: Assignment of the VCD spectrum of the (R)**Ibu** – (S)**Pro** pair.

peak position [cm ⁻¹]	sign	assignment
578	-	Ibu ring deformation (C–H out of plane bending)
642	+	Pro N–H rocking coupled to ring deformations
782	-	Ibu ring deformation coupled to skeletal movements
792	+	Ibu ring deformation coupled to skeletal movements
1032	-	Pro asymmetric C–H bending
1089	+	Ibu O–H wagging coupled to Pro ring deformations
1291	-	Pro *C–H wagging
1396	+	Ibu C–O–H bending
1746	-	Ibu C=O stretching
2889	-	Ibu O–H stretching

 Table TS39: Assignment of the VCD spectrum of the (R)**Nor** – (R)**Pro** pair.

peak position [cm ⁻¹]	sign	assignment
264	-	Nor *C–O–H wagging
272	+	Nor NH ₂ rocking
545	-	Nor ring O–H wagging
589	-	Nor ring O–H wagging
795	+	Nor ring O–H out of plane bending
1129	+	Nor skeletal deformations
1221	-	Nor *C–H wagging
1377	+	Nor NH ₂ wagging
3464	+	Pro N–H stretching coupled to Nor O–H stretching

 Table TS40: Assignment of the VCD spectrum of the (S)**Nor** – (R)**Pro** pair.

peak position [cm ⁻¹]	sign	assignment
287	-	Nor NH ₂ rocking
394	-	Nor ring O–H wagging
804	+	Nor ring C–H out of plane bending
1222	+	Nor *C–H wagging coupled to C–O–H bending
1267	-	Pro C–O–H bending coupled to ring deformations
1607	+	Nor ring C–C stretching

Table TS41: Assignment of the VCD spectrum of the (S)**Bac** – (S)**Pro** pair.

peak position [cm ⁻¹]	sign	assignment
248	+	Bac NH ₂ rocking
1331	-	Pro ring deformations
1342	-	Bac C–O–H bending coupled to Pro *C–H wagging
1354	-	Bac *C–H wagging
1358	+	Bac C–O–H bending coupled to Pro C–O–H bending and Bac *C–H wagging
1756	+	Pro C=O stretching
3405	+	Bac O–H stretching

Table TS42: Assignment of the VCD spectrum of the (R)**Bac** – (S)**Pro** pair.

peak position [cm ⁻¹]	sign	assignment
262	-	Bac NH ₂ rocking
534	-	Bac ring deformation (C–H out of plane bending)
589	+	Bac ring deformation coupled to CH ₂ wagging and Pro O–H wagging
1027	-	Bac O–H wagging coupled to Pro ring CH ₂ rocking
1120	-	Pro *C–H wagging coupled to O–H wagging
1263	-	Bac *C–H wagging coupled to CH ₂ rocking
1743	-	Bac C=O stretching
2987	+	Bac O–H stretching coupled to Pro C–H stretching of CH ₂ group in α position to N–H group
3000	-	Bac O–H stretching coupled Pro ring C–H stretching
3009	-	Bac O–H stretching coupled Pro ring C–H stretching

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