

Analyzing the symmetrical arrangement of structural repeats in proteins with CE-Symm [S1 Tab]

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Type	Count	Percentage
Asymmetric	747	74.2%
Rotational	214	21.2%
C2	160	74.8%
C3	10	4.7%
C4	2	0.9%
C5	3	1.4%
C6	9	4.2%
C7	10	4.7%
C8	20	9.3%
Dihedral	18	1.8%
D2	14	77.8%
D3	1	5.6%
D4	2	11.1%
D5	1	5.6%
Helical	11	1.1%
H2	9	81.8%
H3	2	18.2%
H10	1	9.1%
Superhelical	2	0.2%
Repeats	15	1.5%

S1 Tab. Summary of the updated annotations in the internal symmetry benchmarking dataset.