

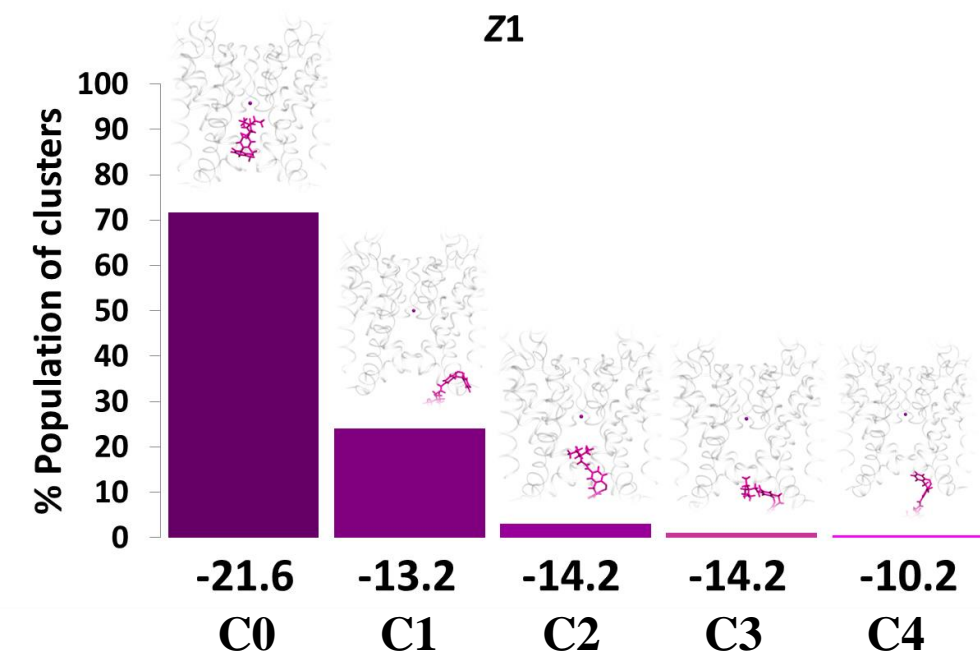
Representative structures from Hierarchical clustering result

Deciphering the Shape Selective Conformational Equilibrium of *E*- and *Z*-Locked Azobenzene-Tetraethyl Ammonium Ion in Regulating Photo-switchable K⁺-ion Channel Blocking

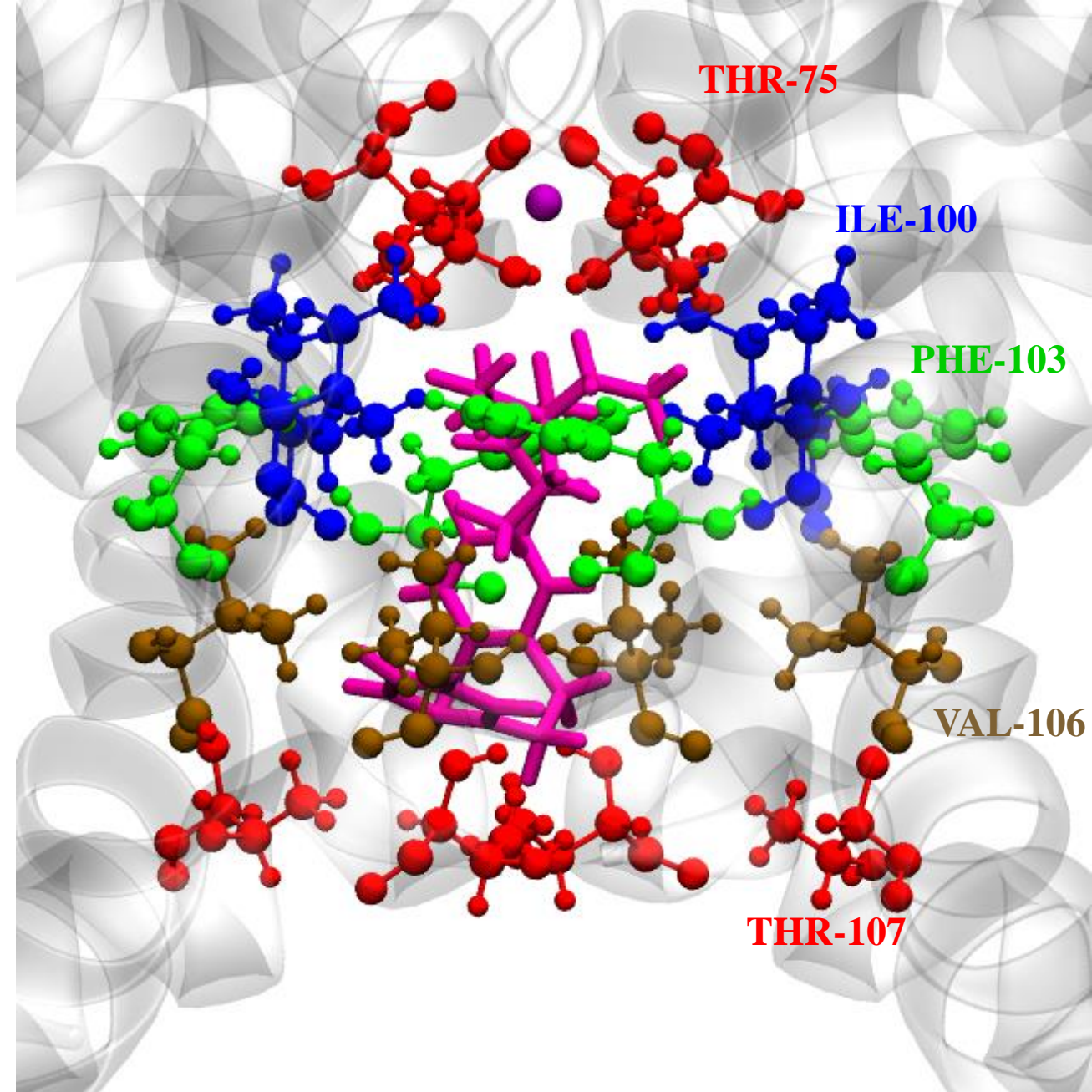
Rinsha Cholasseri^a and Susmita De^{*b}

^aTheoretical and Computational Chemistry Laboratory, Department of Chemistry, National Institute of Technology Calicut, Kozhikode, Kerala, India - 673 601.

^bDepartment of Chemistry, University of Calicut, Calicut University P. O., Malappuram, Kerala, India - 673 635. E-mail: susmita@uoc.ac.in

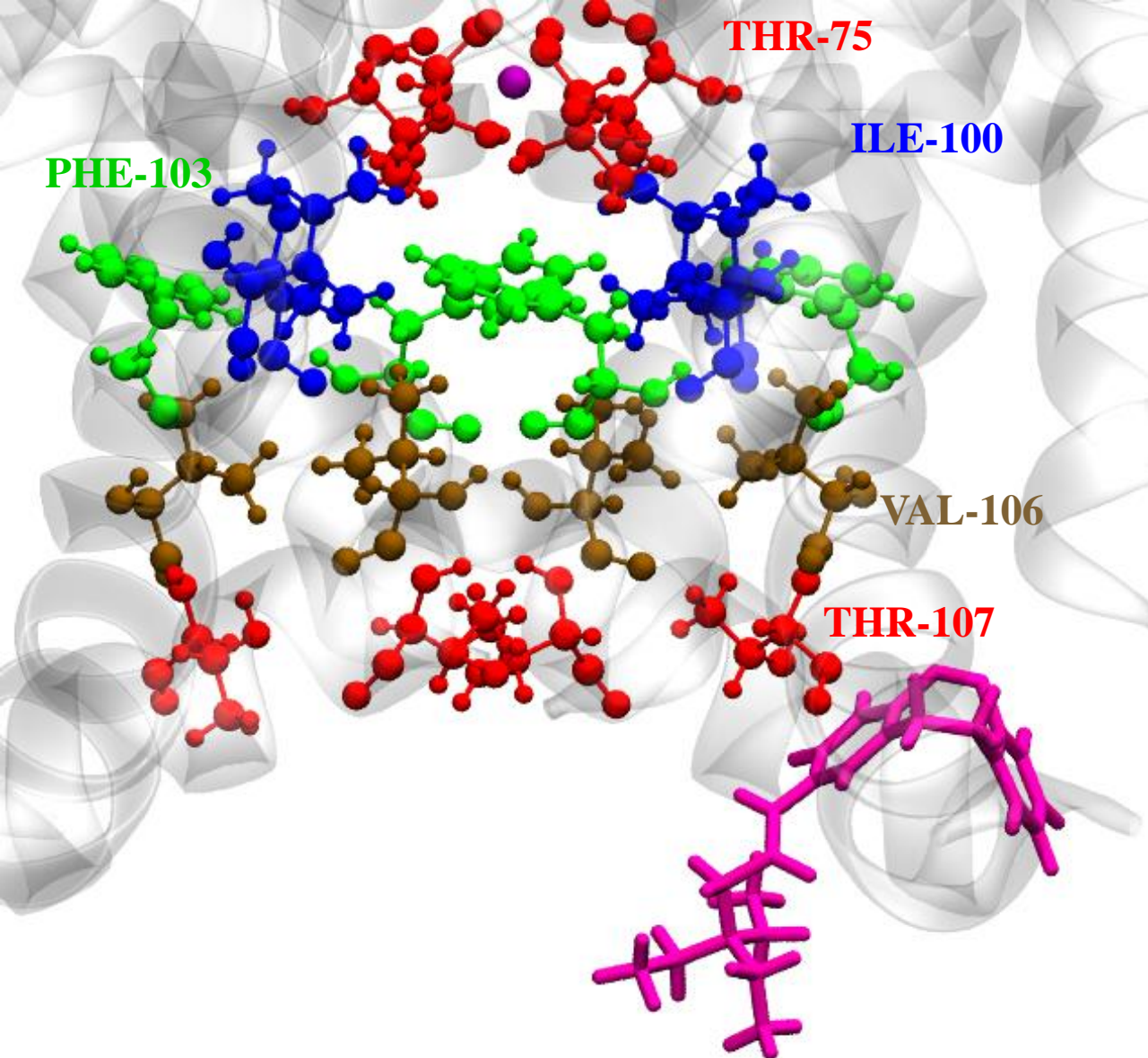


Cn (n = 0 – 1) indicate different clusters. The percentage population of clusters is given on the Y-axis. Corresponding binding energy (B.E.) is given in kcal/mol on the X-axis. Protein is represented in new cartoon model with K⁺-ion (purple ball), and Z-LAB-TEA is shown in stick model (magenta colour).



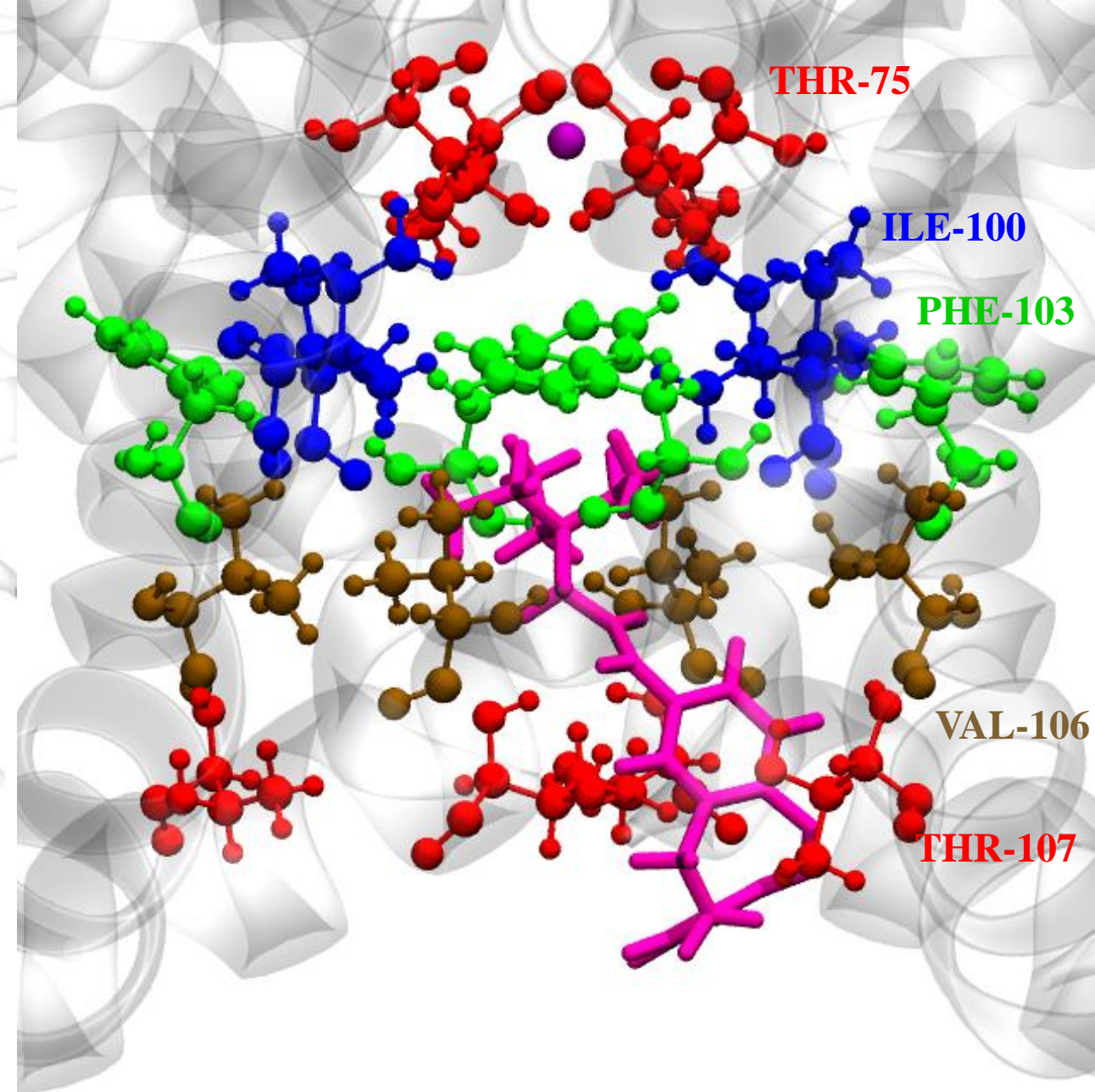
Z1(C0)

MM/PBSA Binding energy = -21.6 kcal/mol²



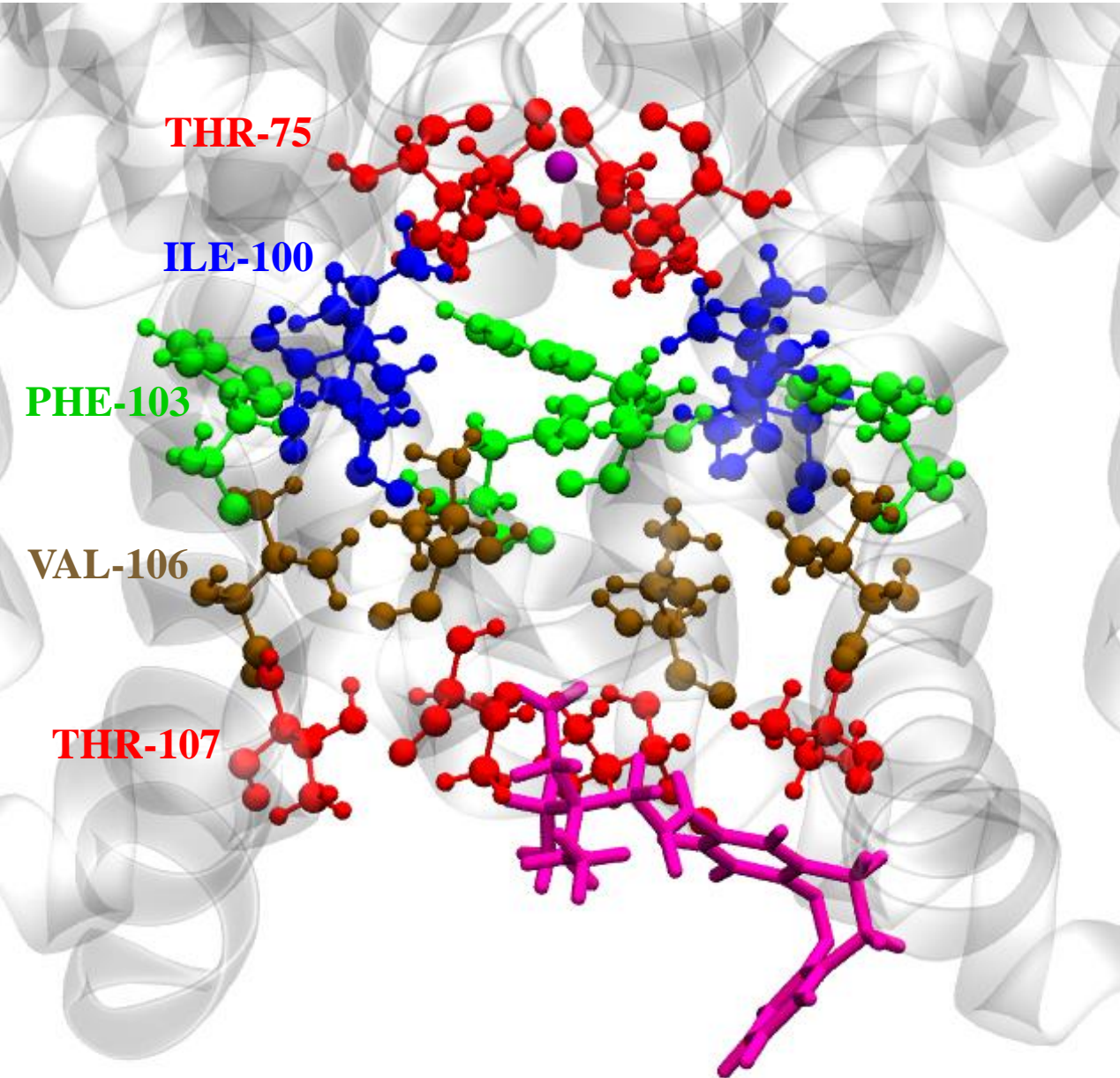
Z1 (C1)

MM/PBSA Binding energy = -13.2 kcal/mol



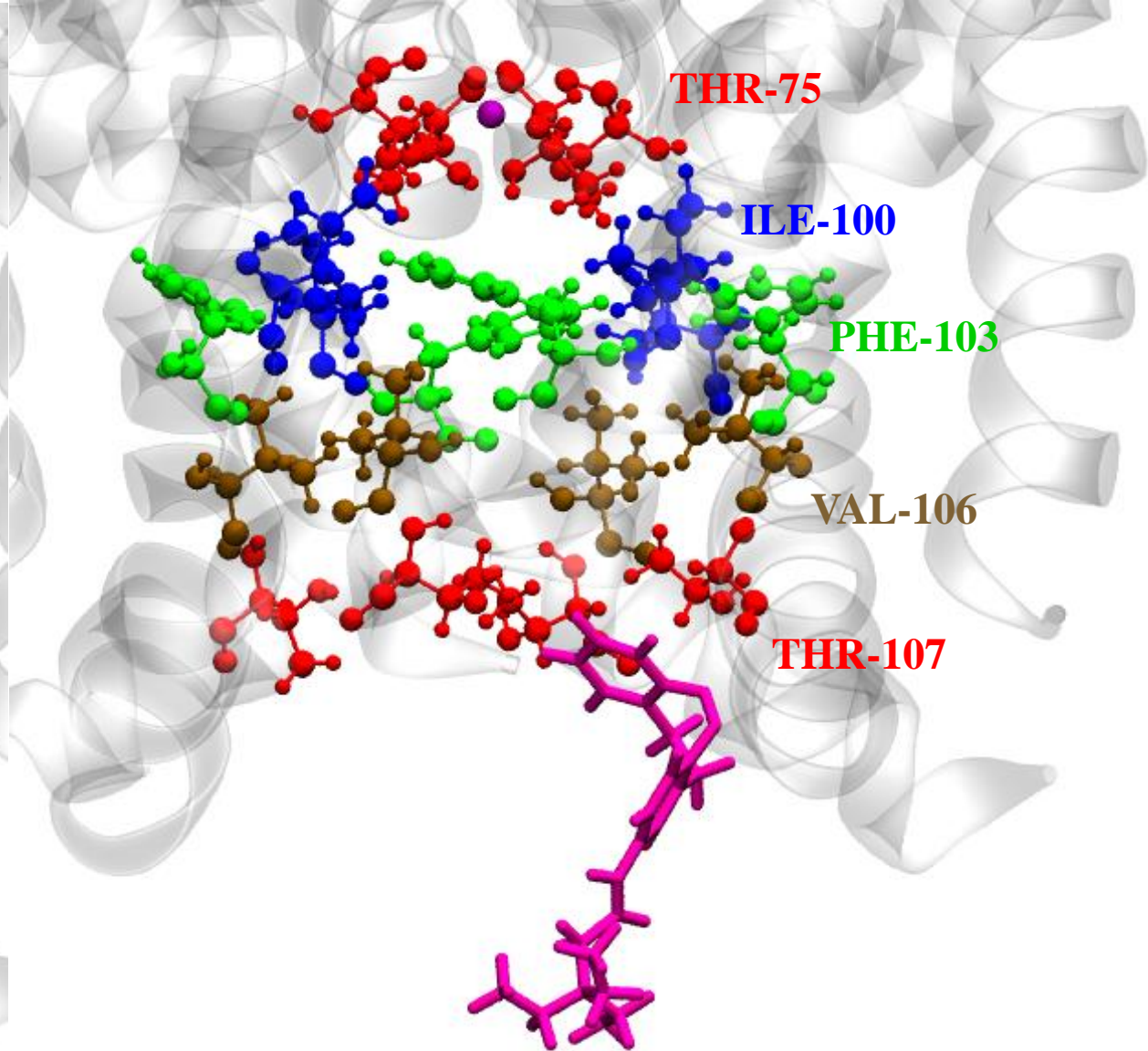
Z1 (C2)

MM/PBSA Binding energy = -14.2 kcal/mol³



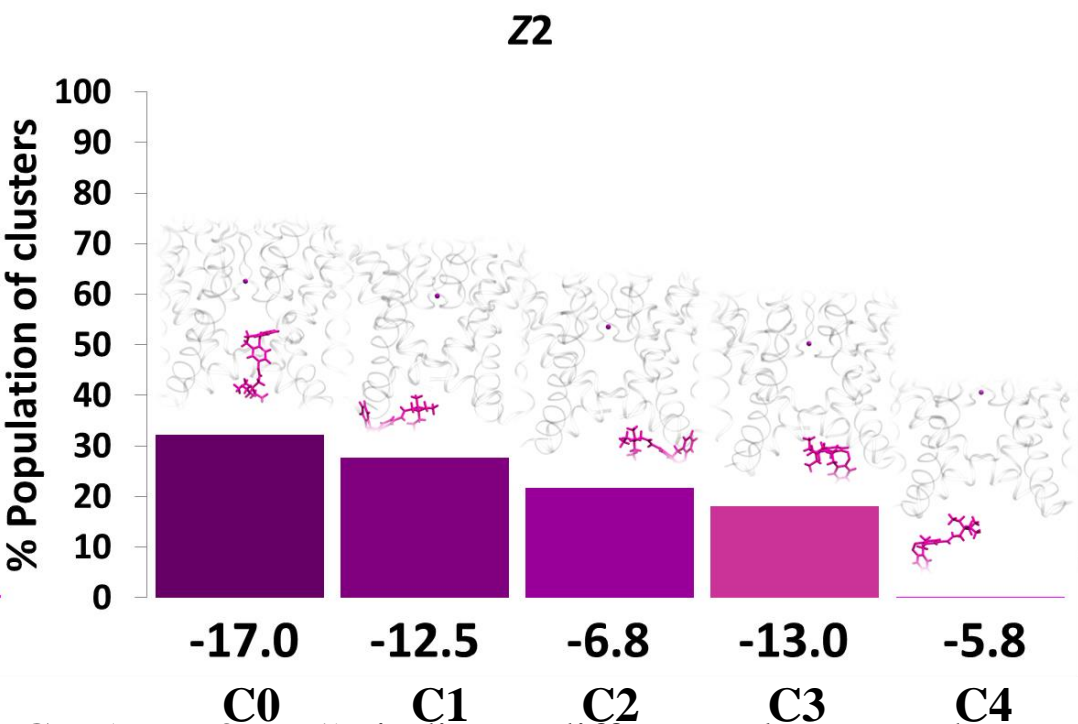
Z1 (C3)

MM/PBSA Binding energy = -14.2 kcal/mol

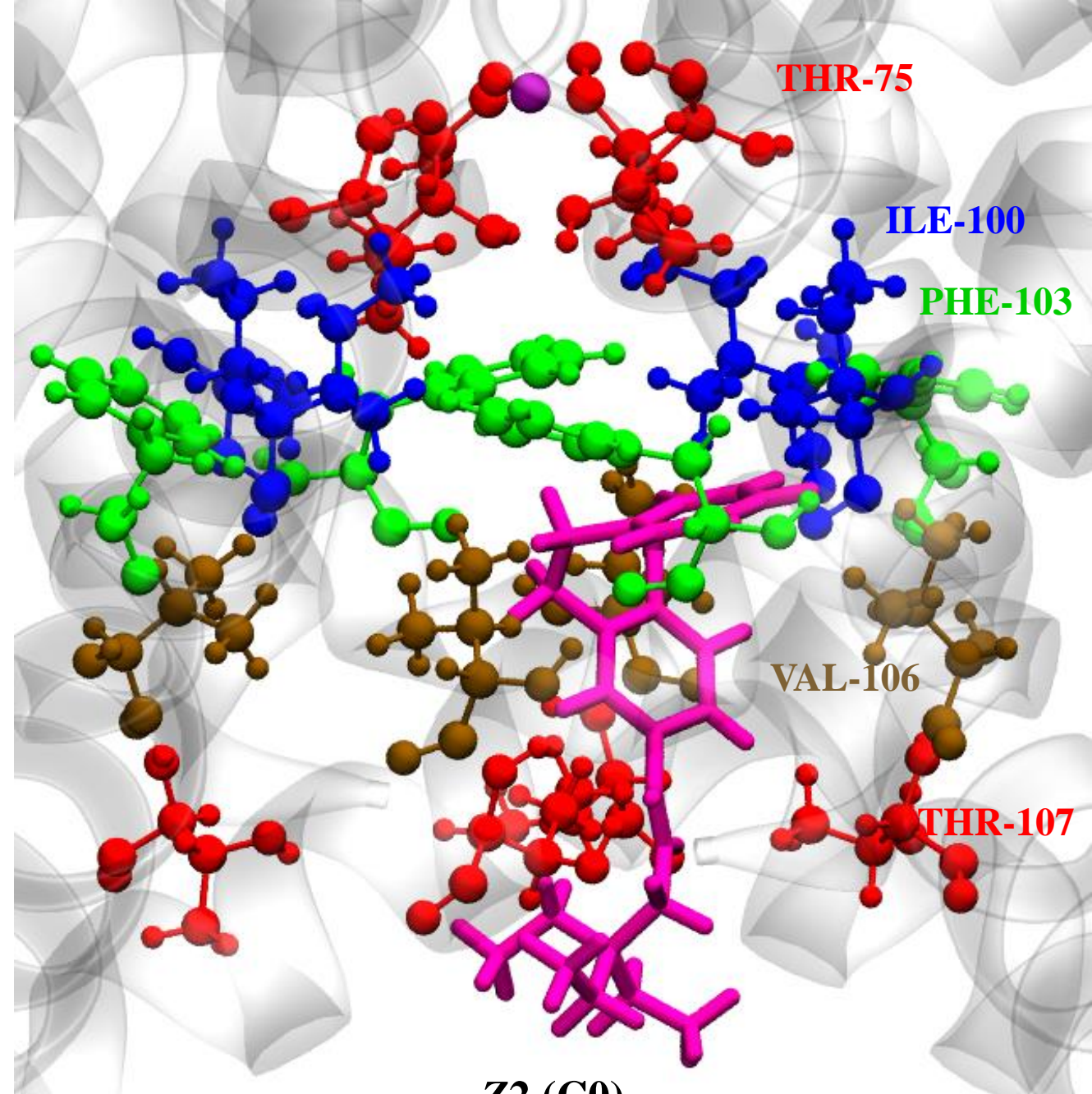


Z1 (C4)

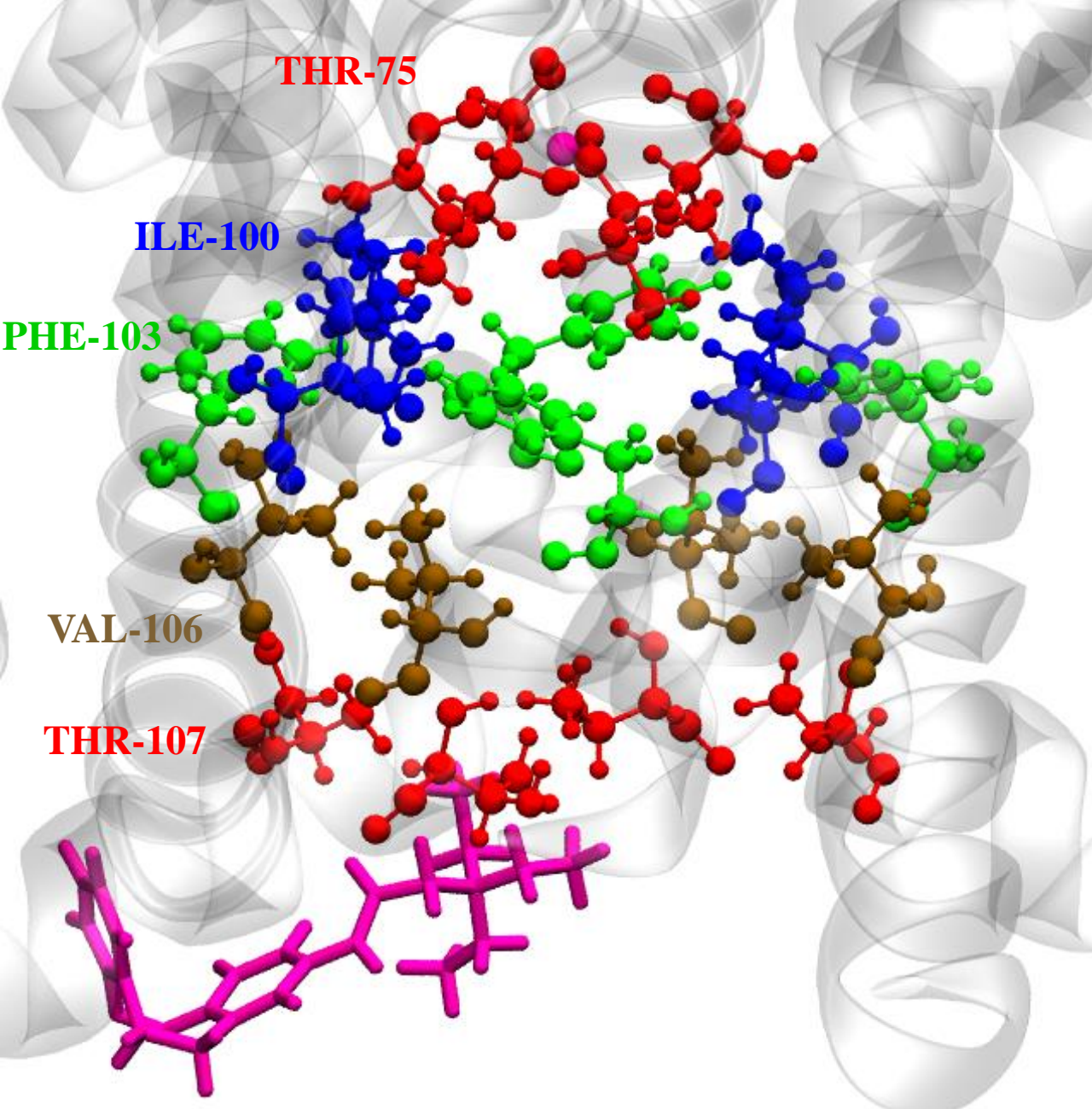
MM/PBSA Binding energy = -10.2 kcal/mol⁴



C_n ($n = 0 - 1$) indicate different clusters. The percentage population of clusters is given on the Y-axis. Corresponding binding energy (B.E.) is given in kcal/mol on the X-axis. Protein is represented in new cartoon model with K^+ -ion (purple ball), and Z-LAB-TEA is shown in stick model (magenta colour).

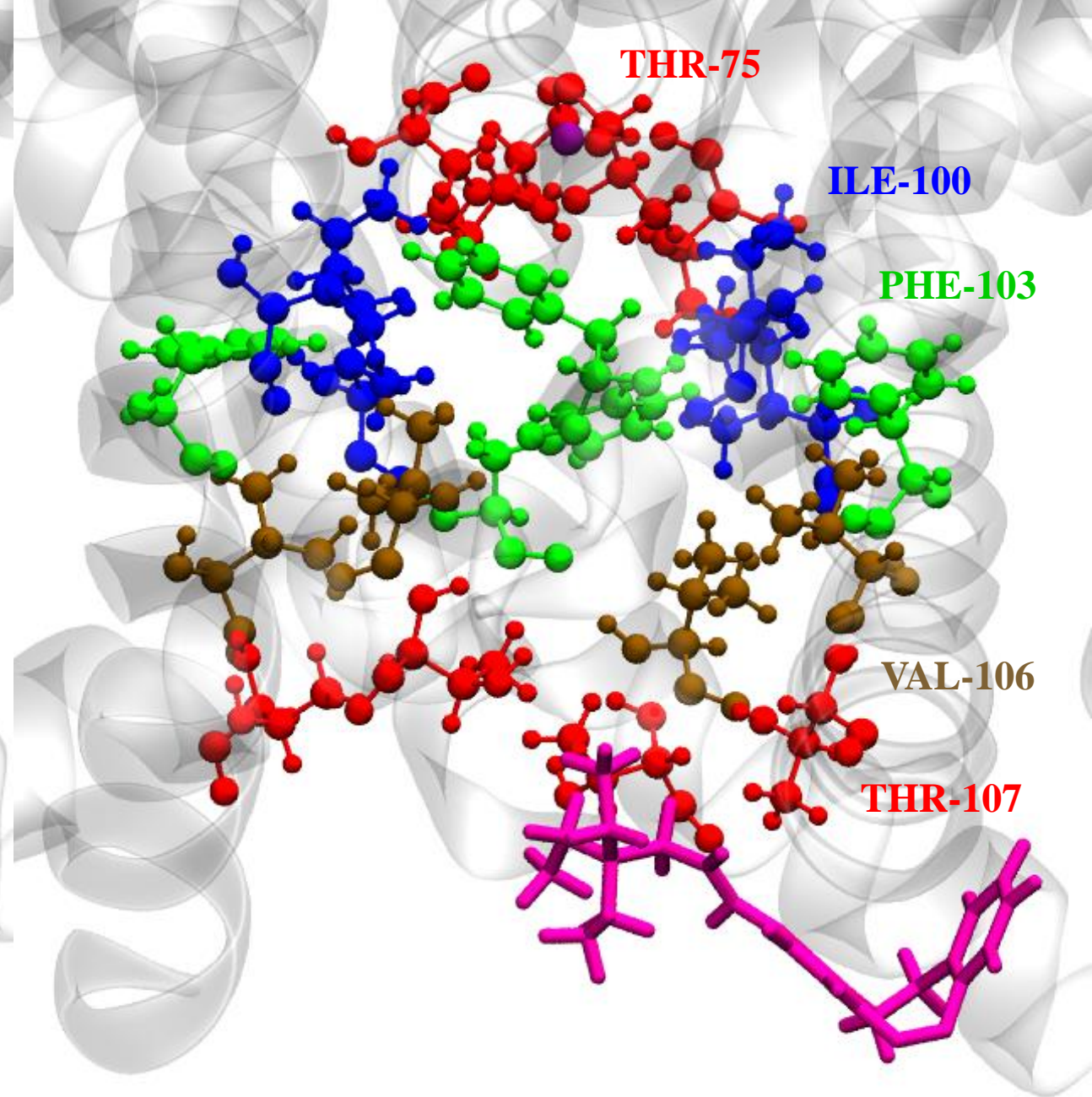


MM/PBSA Binding energy = -17.0 kcal/mol⁵



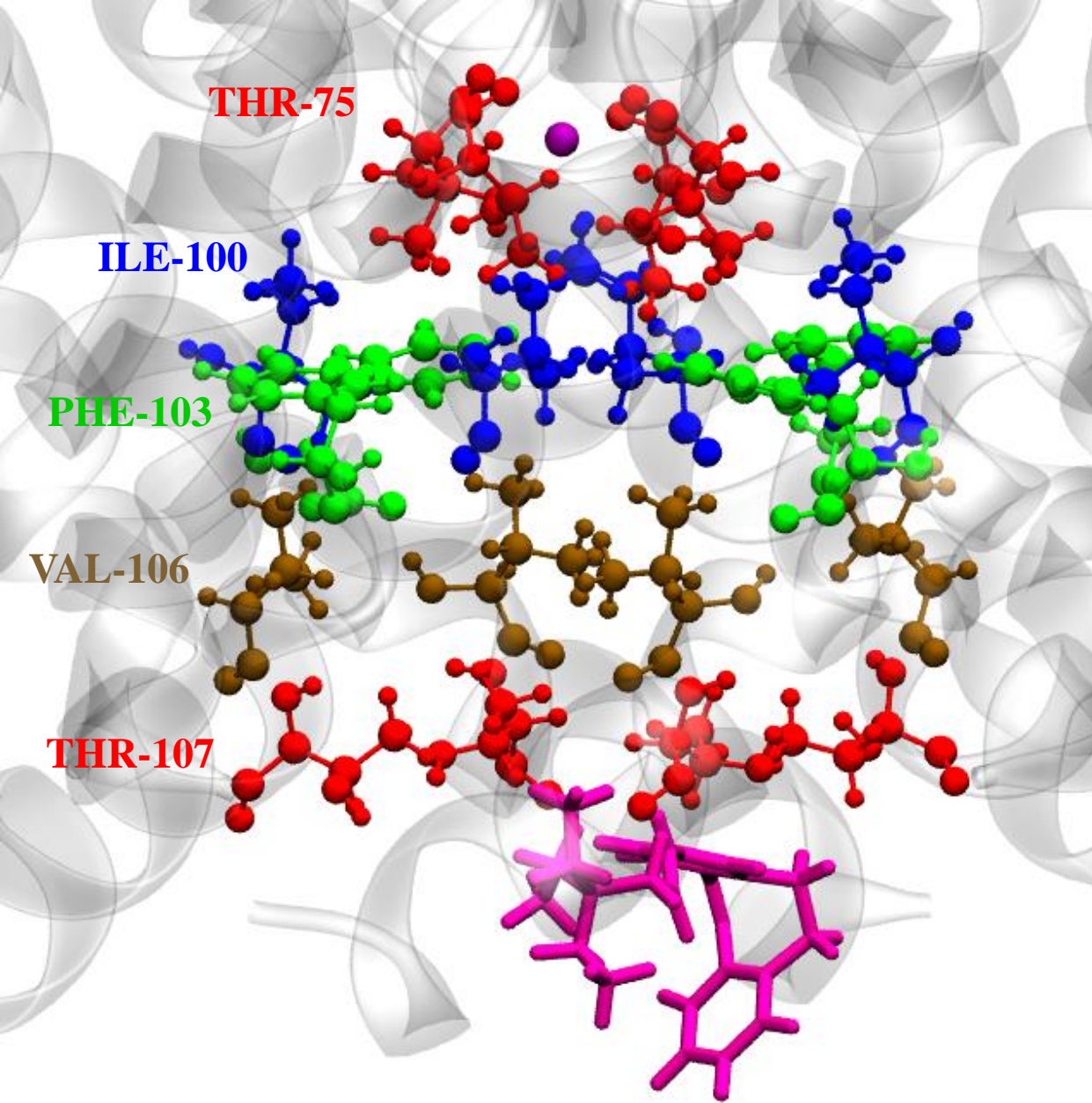
Z2 (C1)

MM/PBSA Binding energy = -12.5 kcal/mol



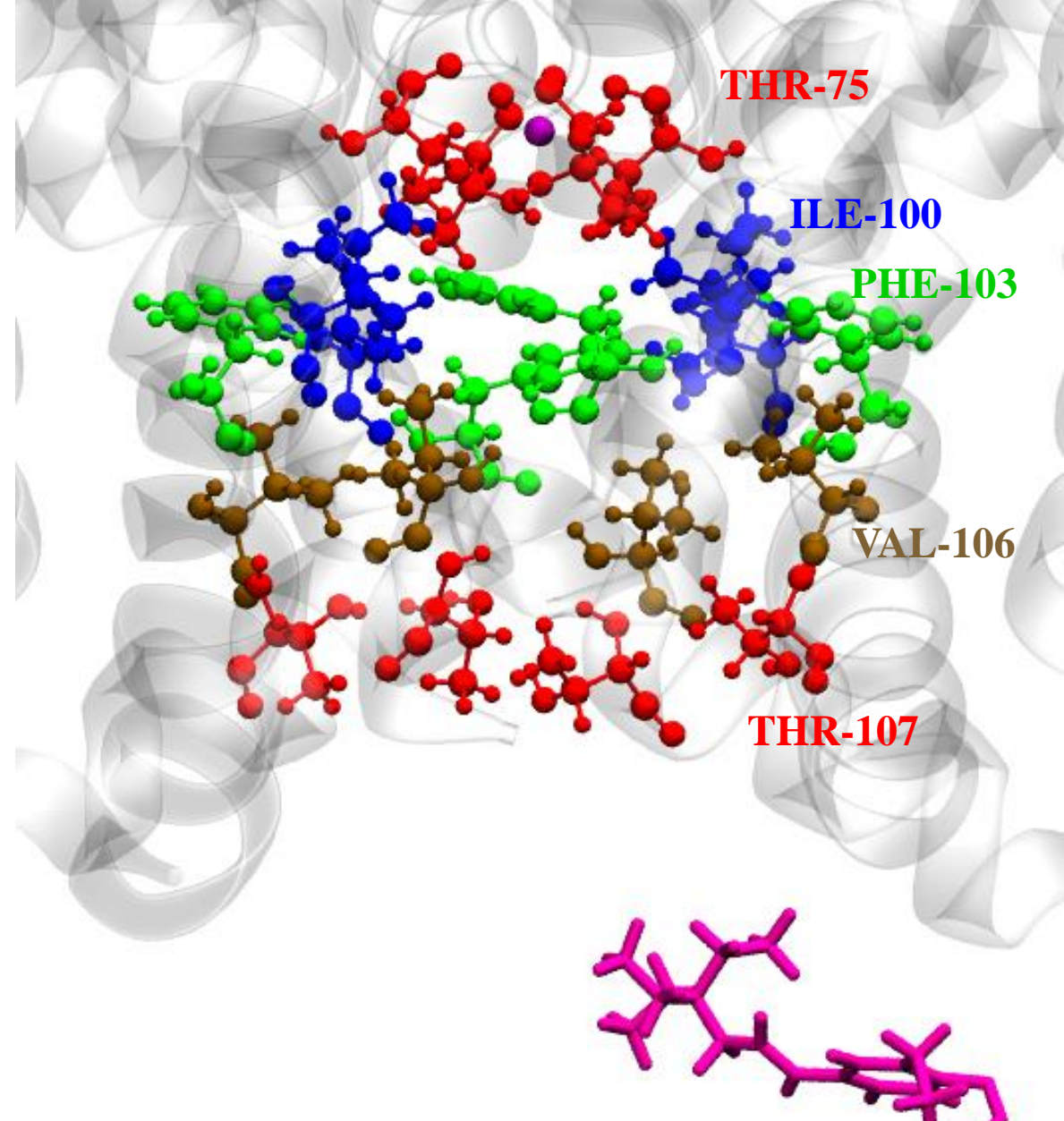
Z2 (C2)

MM/PBSA Binding energy = -6.8 kcal/mol



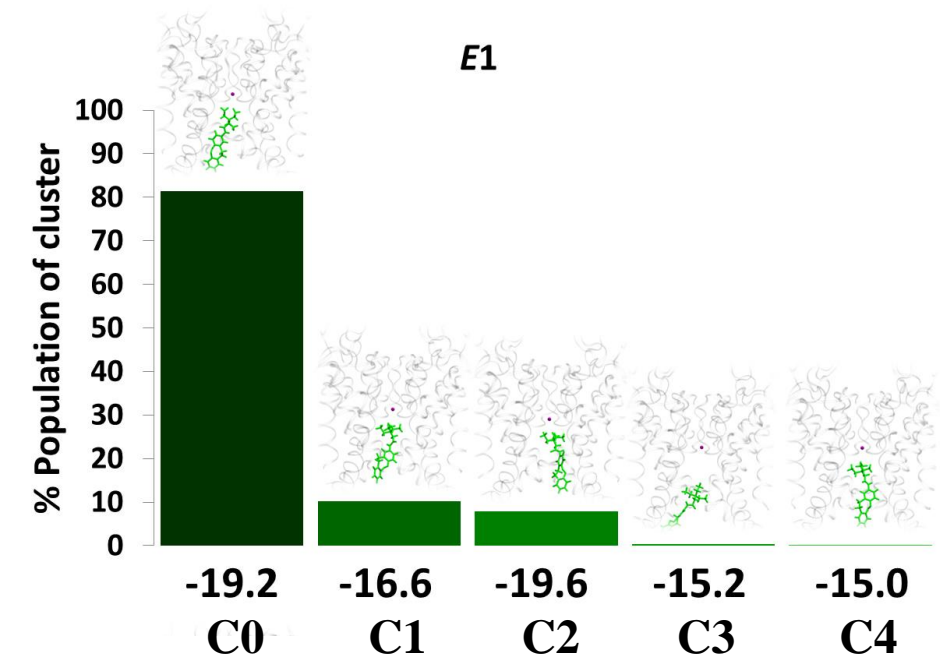
Z2 (C3)

MM/PBSA Binding energy = -13.0 kcal/mol

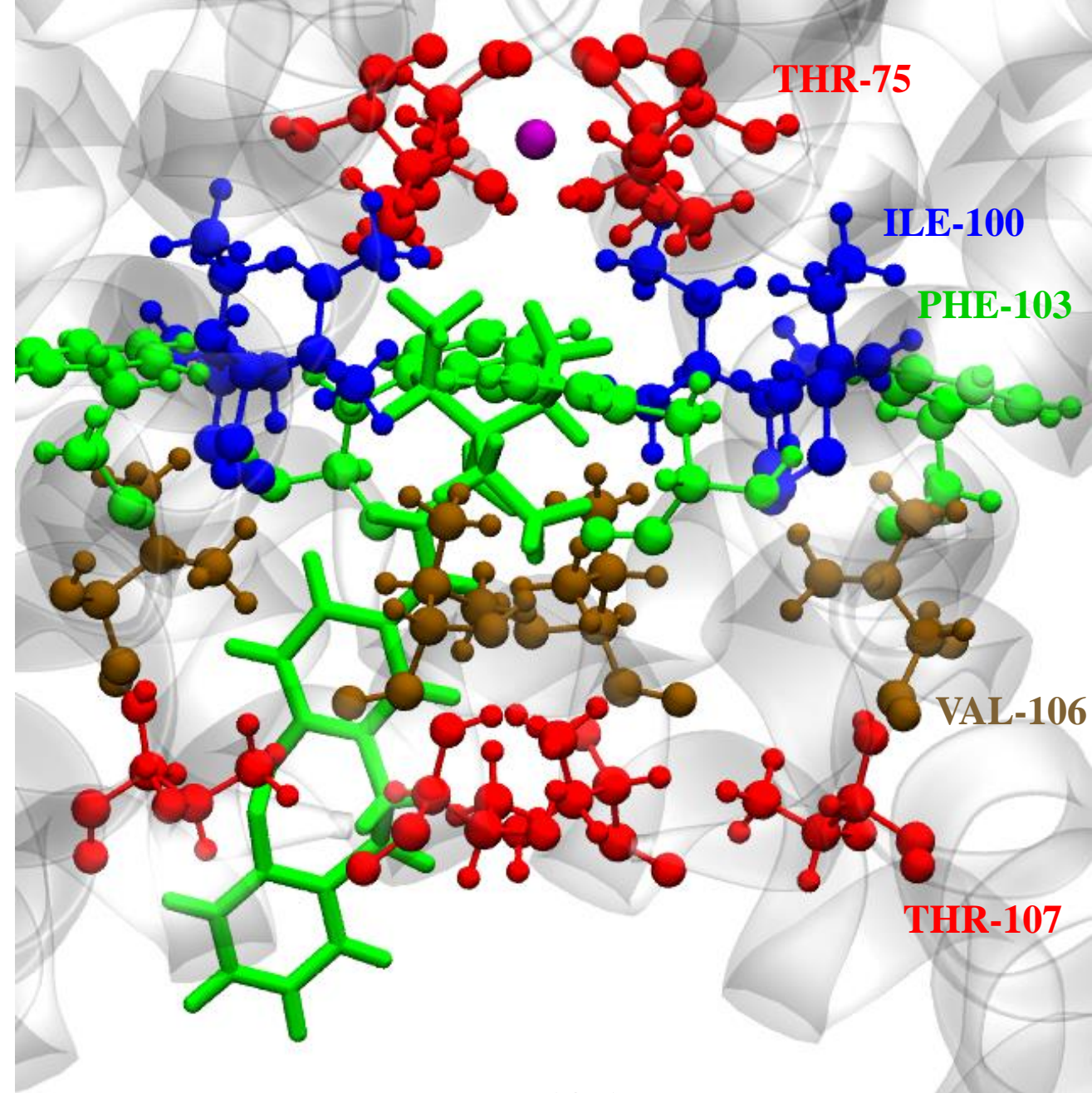


Z2 (C4)

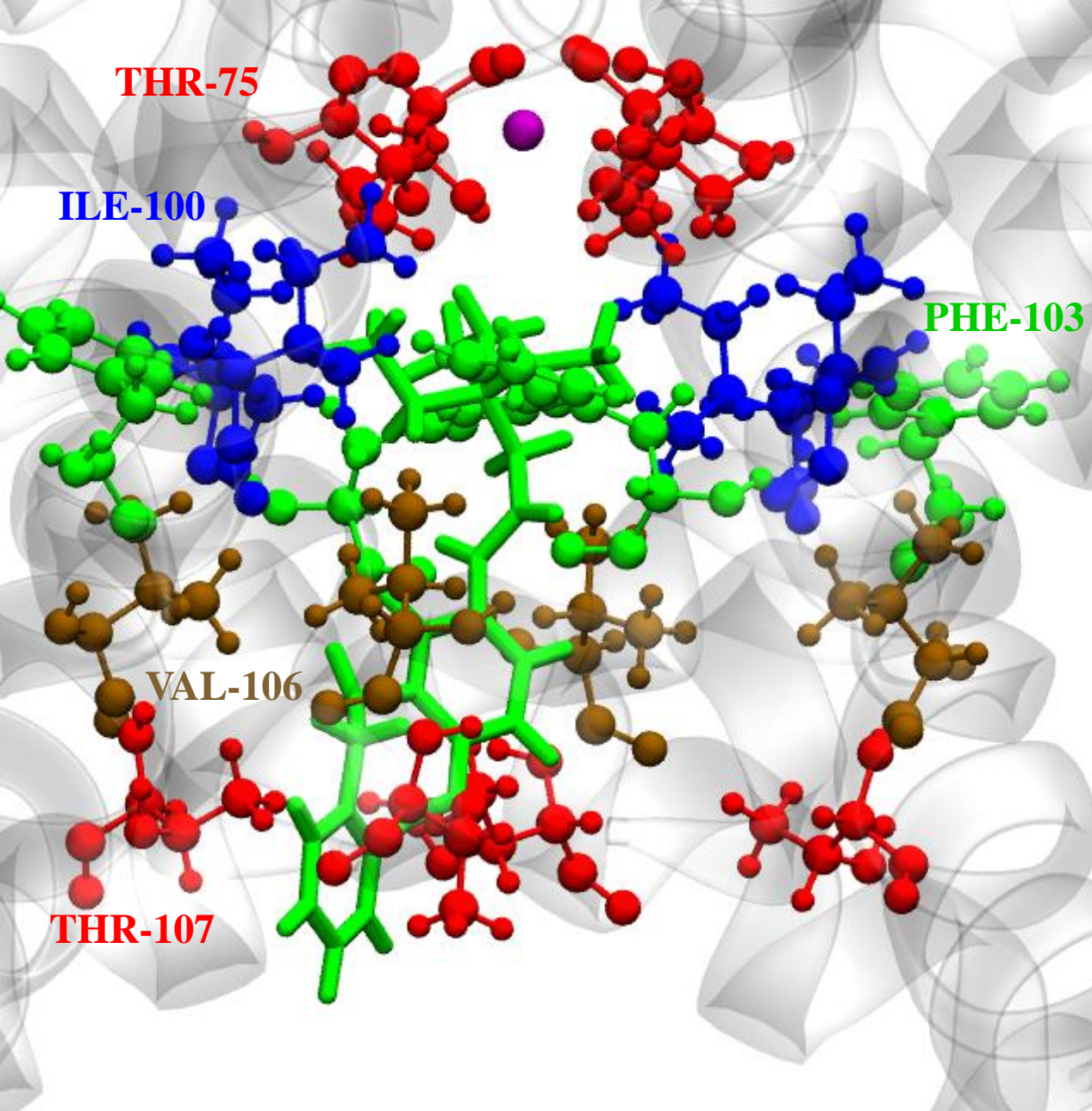
MM/PBSA Binding energy = -5.8 kcal/mol



C_n (n = 0 – 1) indicate different clusters. The percentage population of clusters is given on the Y-axis. Corresponding binding energy (B.E.) is given in kcal/mol on the X-axis. Protein is represented in new cartoon model with K⁺-ion (purple ball), and *E*-LAB-TEA is shown in stick model (green colour).

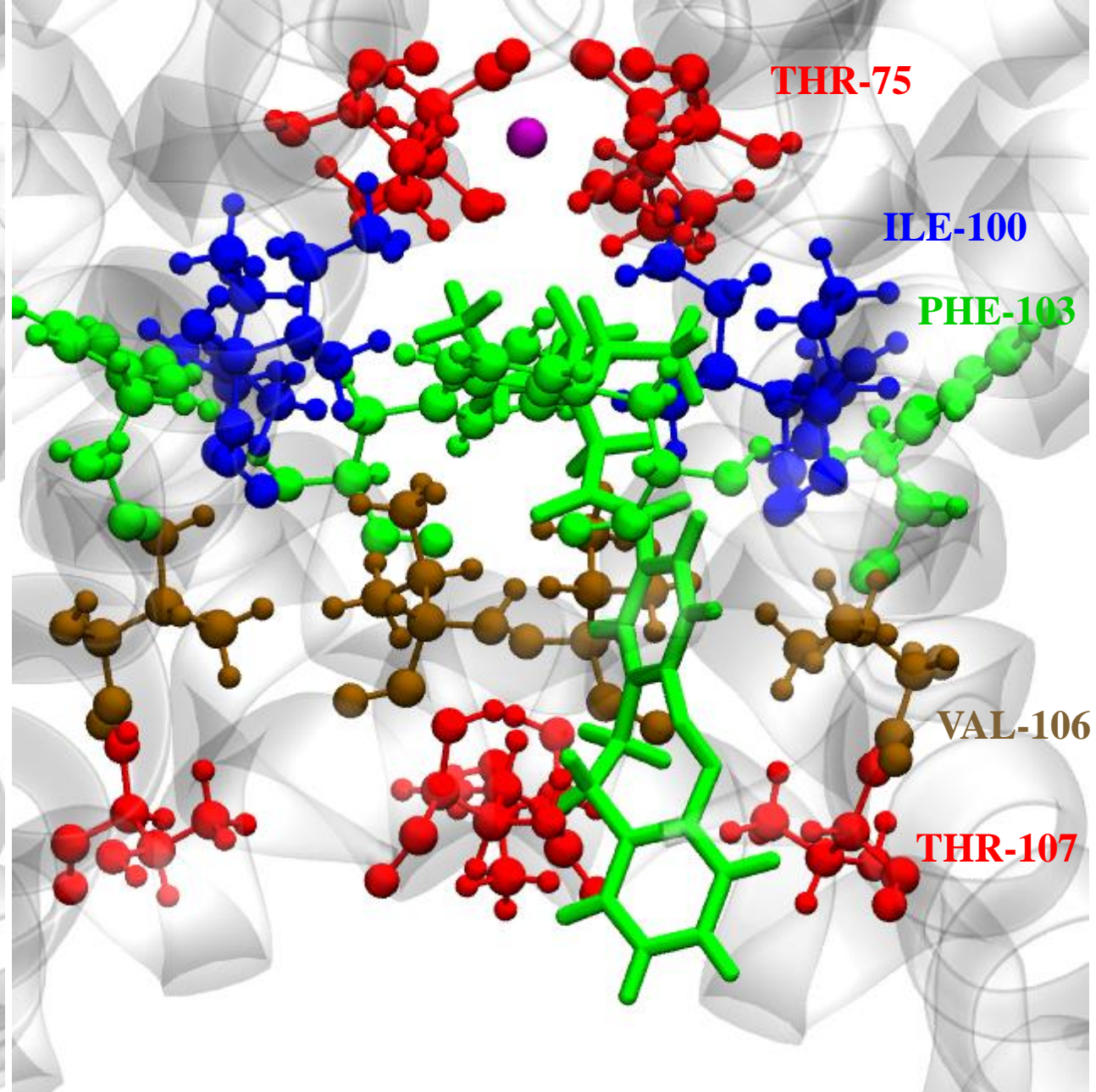


E1 (C0)
MM/PBSA Binding energy = -19.2 kcal/mol⁸



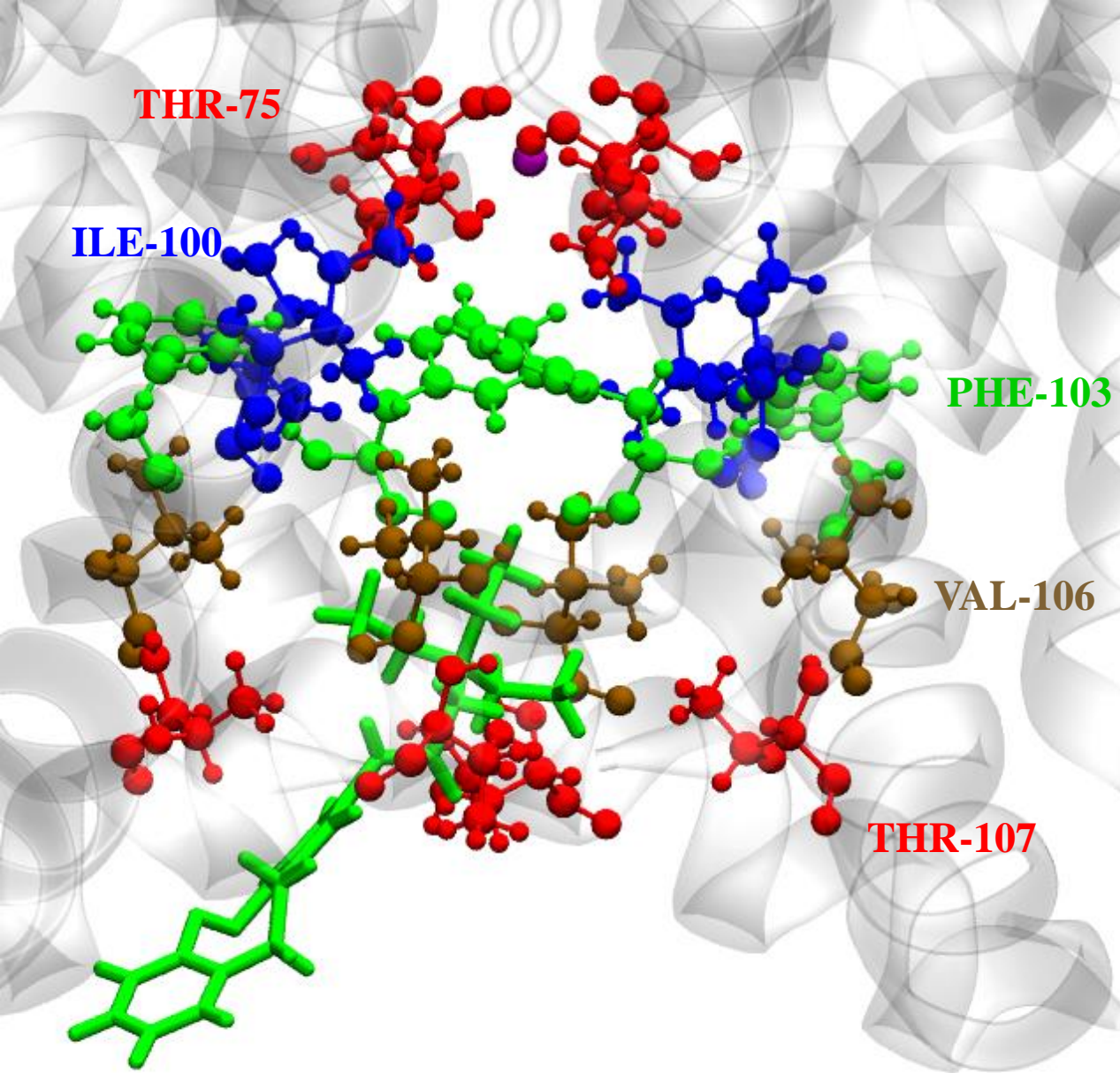
E1 (C1)

MM/PBSA Binding energy = -16.6 kcal/mol



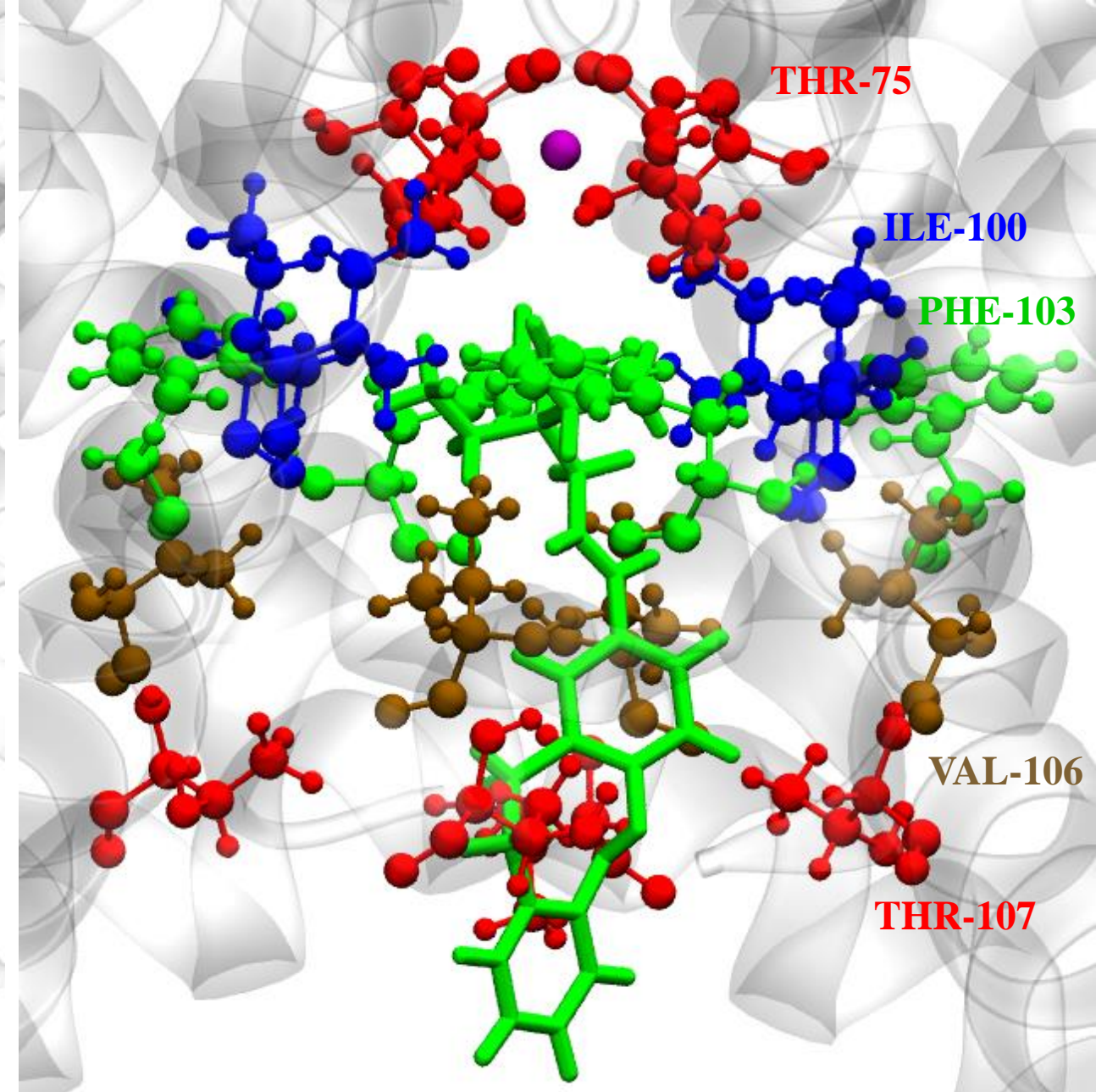
E1 (C2)

MM/PBSA Binding energy = -19.6 kcal/mol⁹



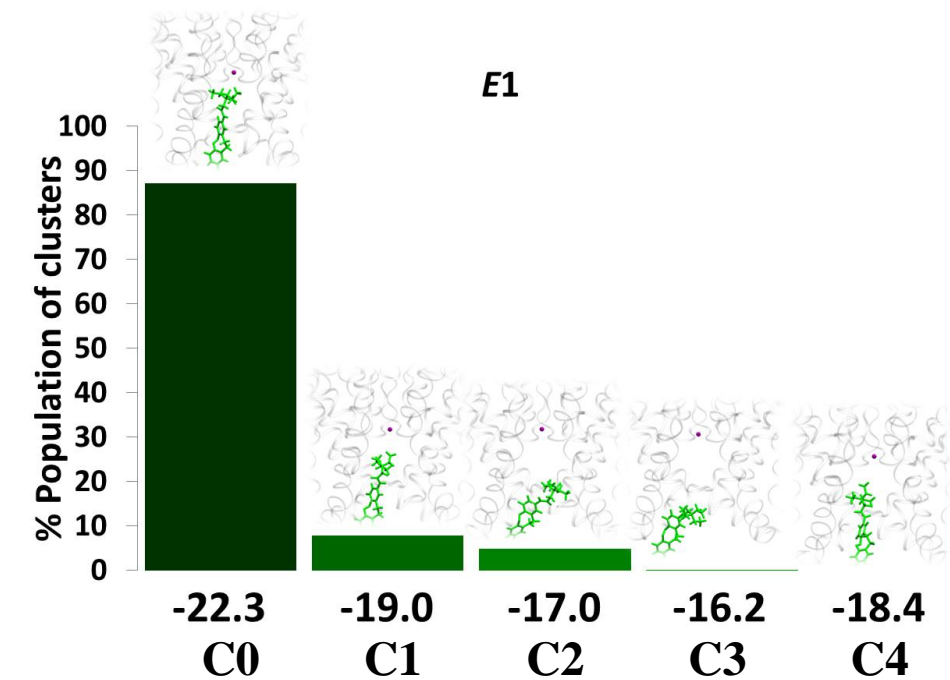
E1 (C3)

MM/PBSA Binding energy = -15.2 kcal/mol

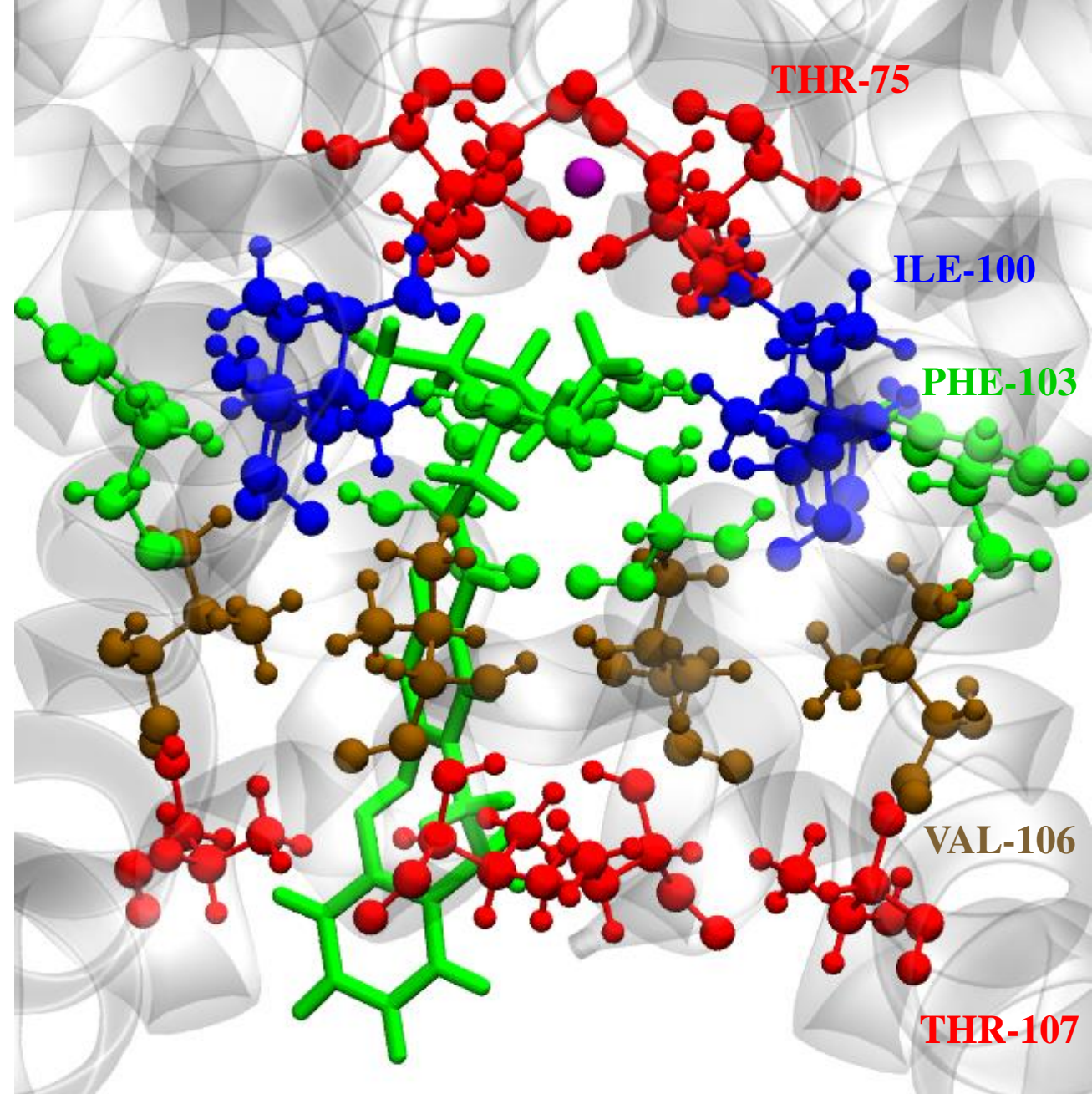


E1 (C4)

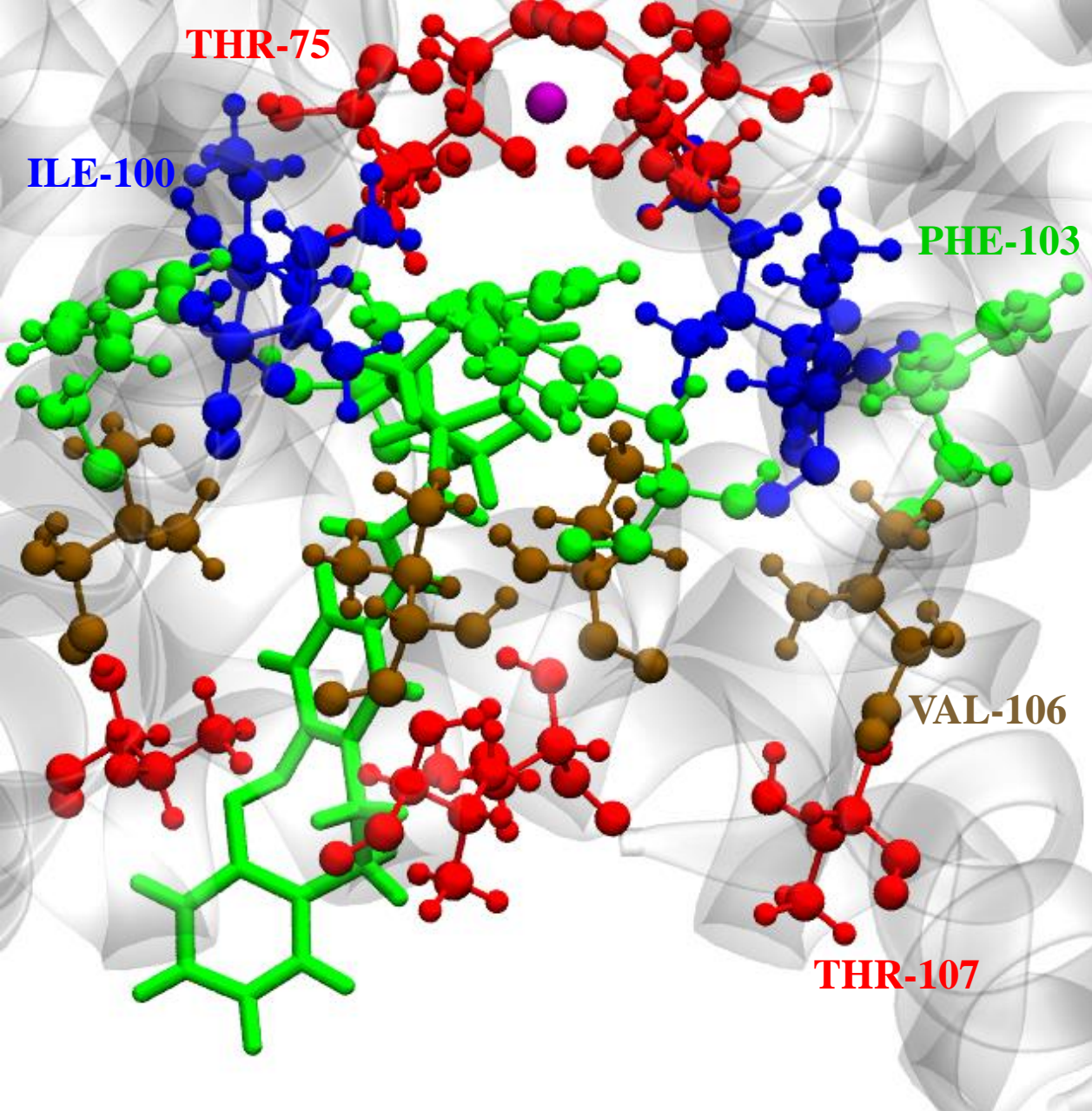
MM/PBSA Binding energy = -15.0 kcal/mol¹⁰



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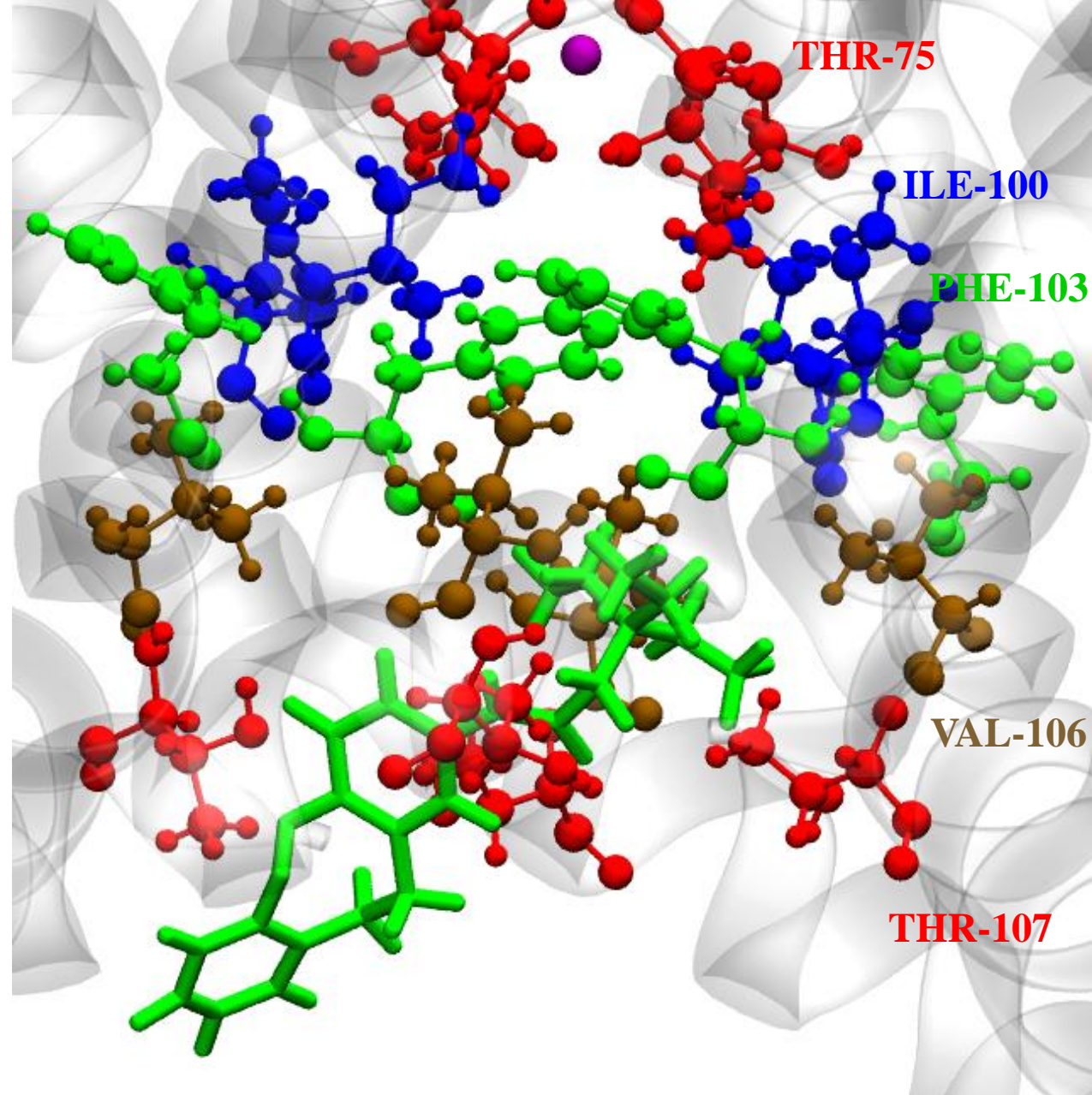


*E*1 (C0)
MM/PBSA Binding energy = -22.3 kcal/mol¹¹



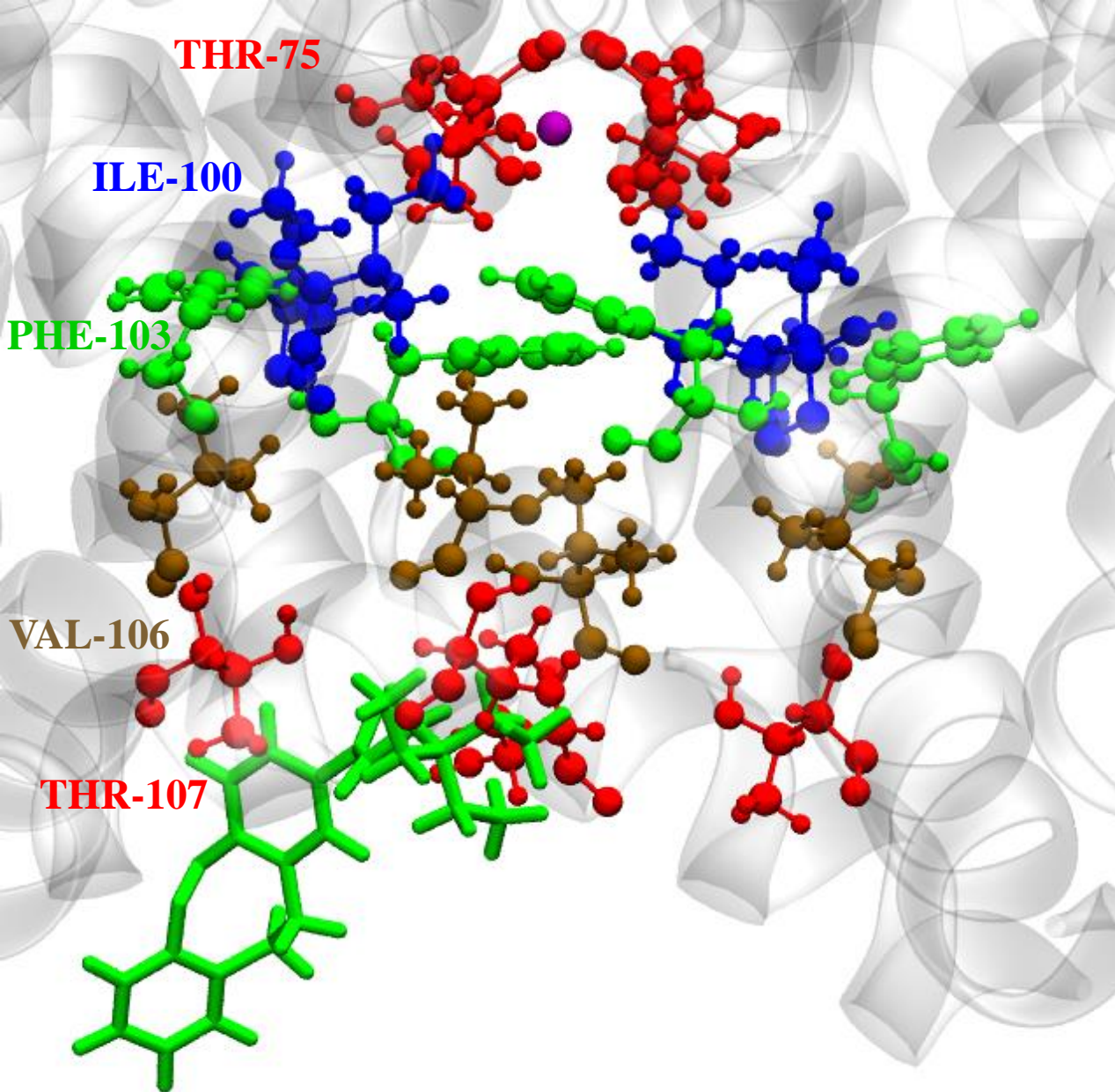
E1 (C1)

MM/PBSA Binding energy = -19.0 kcal/mol



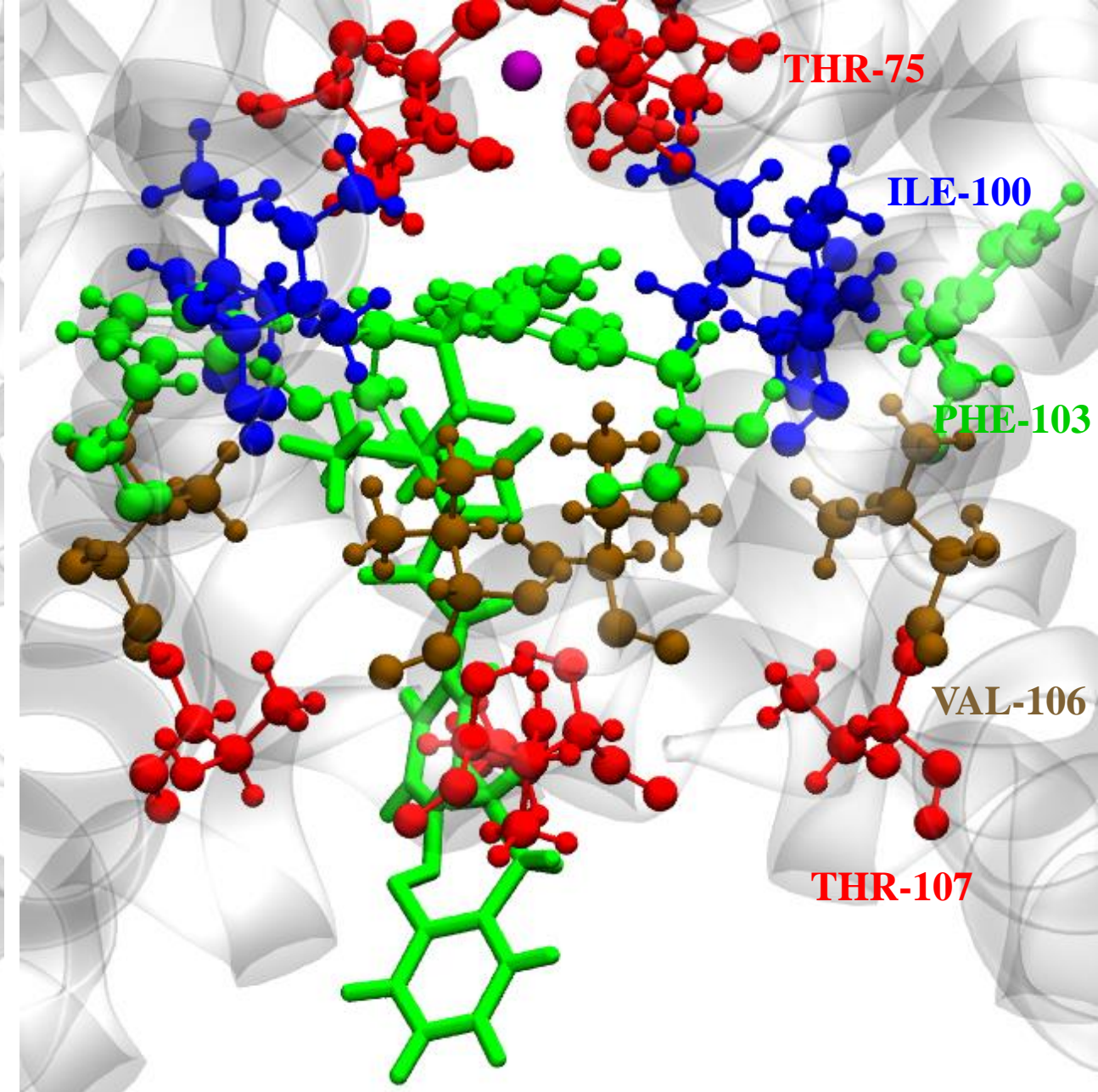
E1 (C2)

MM/PBSA Binding energy = -17.0 kcal/mol¹²



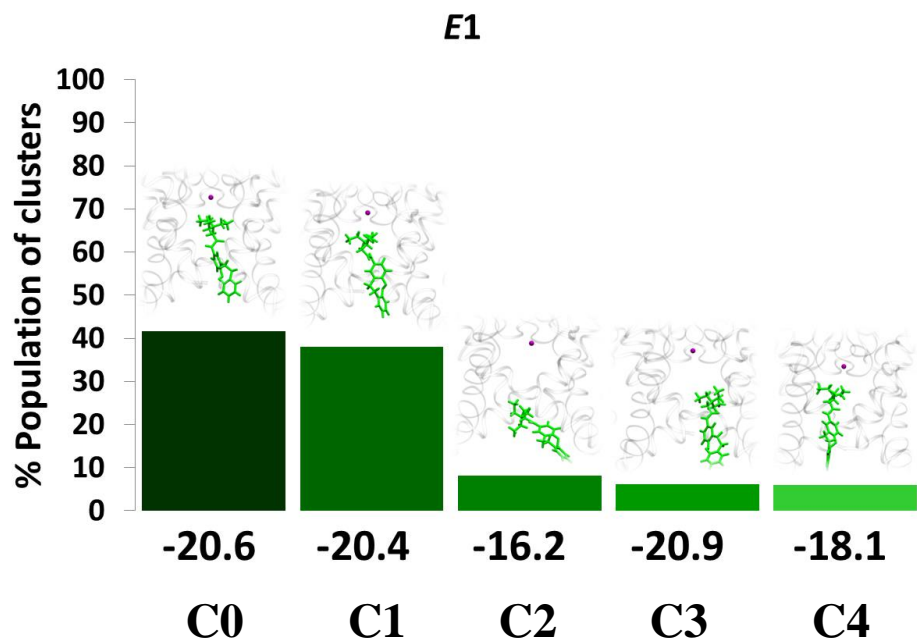
E1 (C3)

MM/PBSA Binding energy = -16.2 kcal/mol

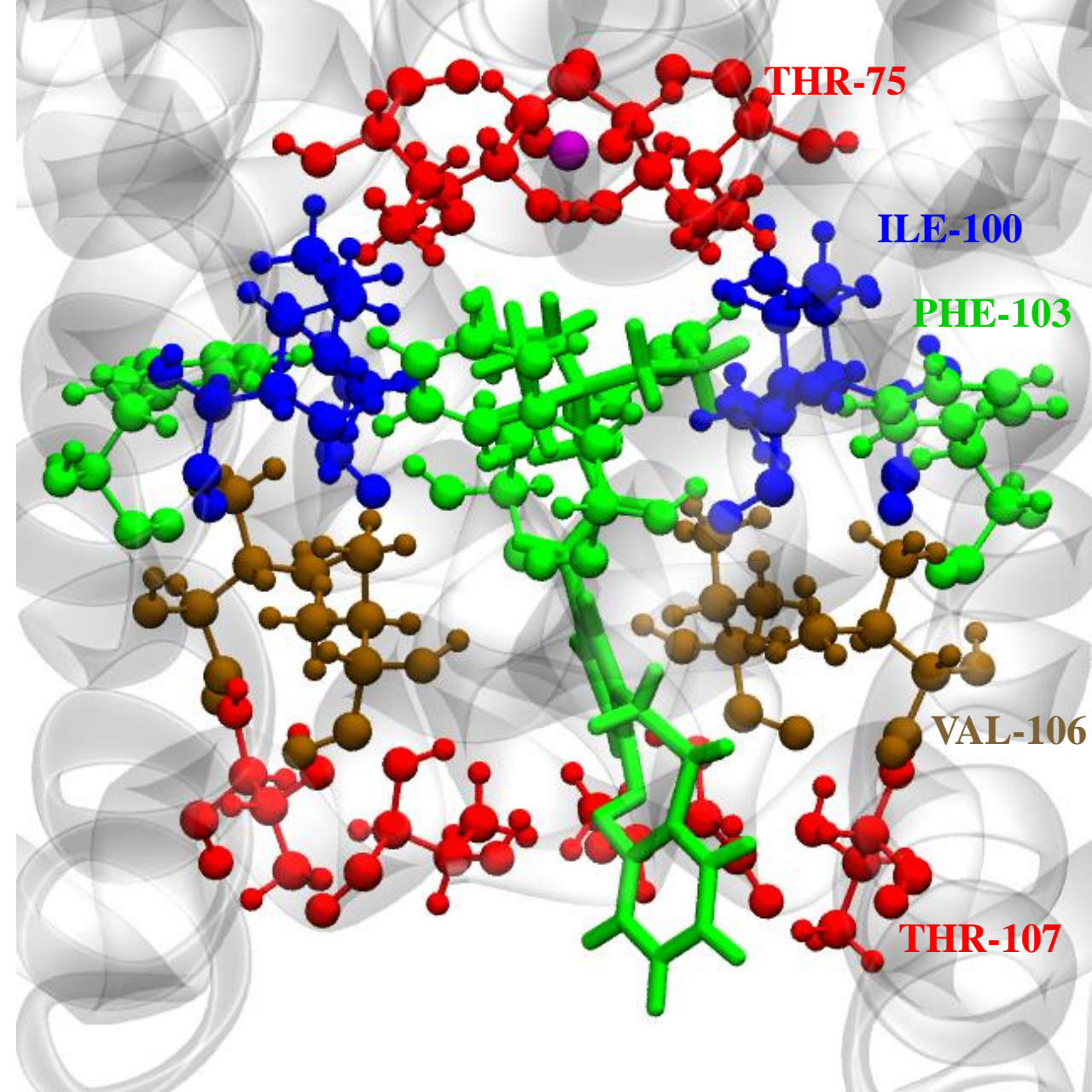


E1 (C4)

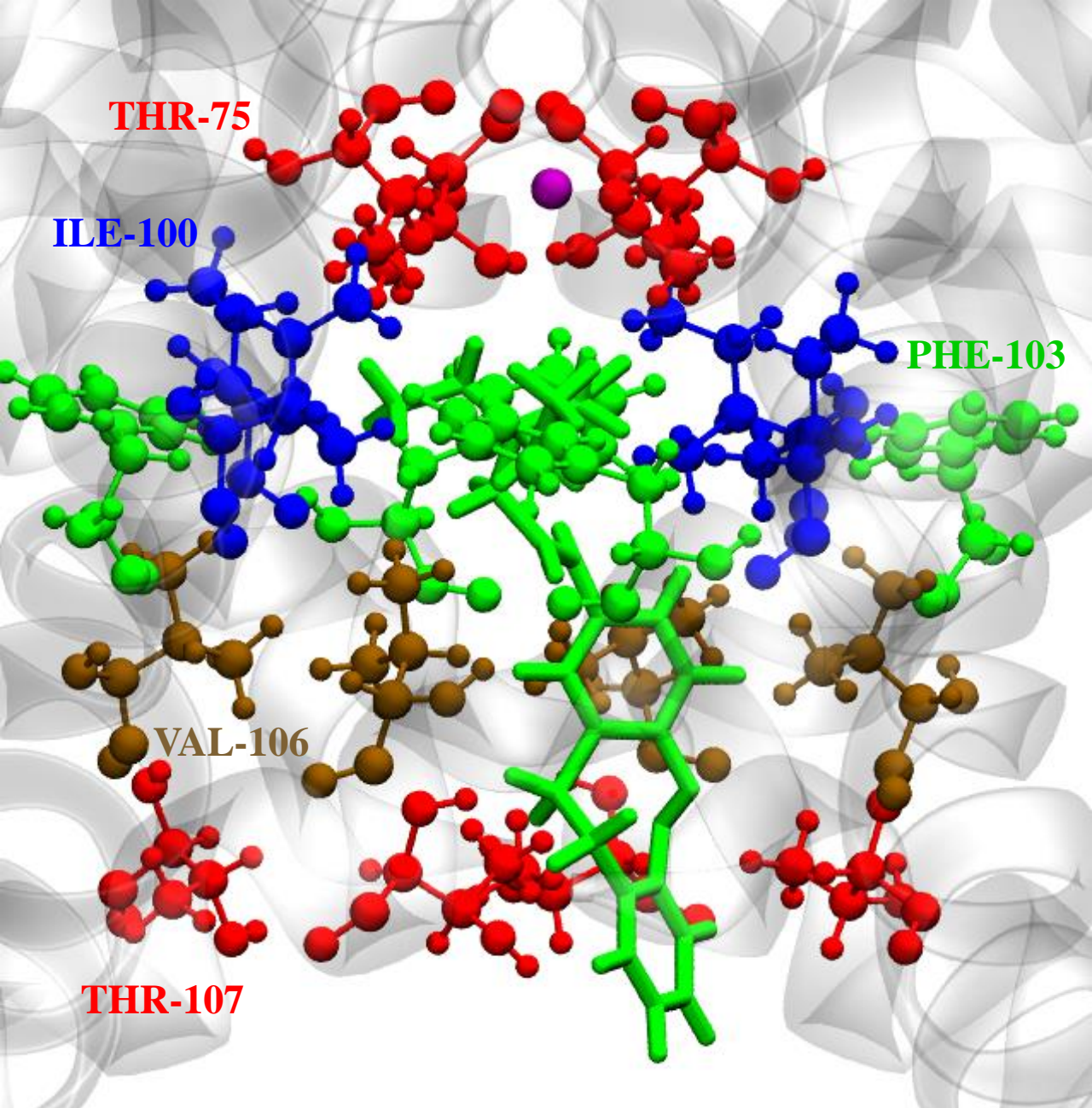
MM/PBSA Binding energy = -18.4 kcal/mol¹³



C_n (n = 0 – 1) indicate different clusters. The percentage population of clusters is given on the Y-axis. Corresponding binding energy (B.E.) is given in kcal/mol on the X-axis. Protein is represented in new cartoon model with K⁺-ion (purple ball), and *E*-LAB-TEA is shown in stick model (green colour).

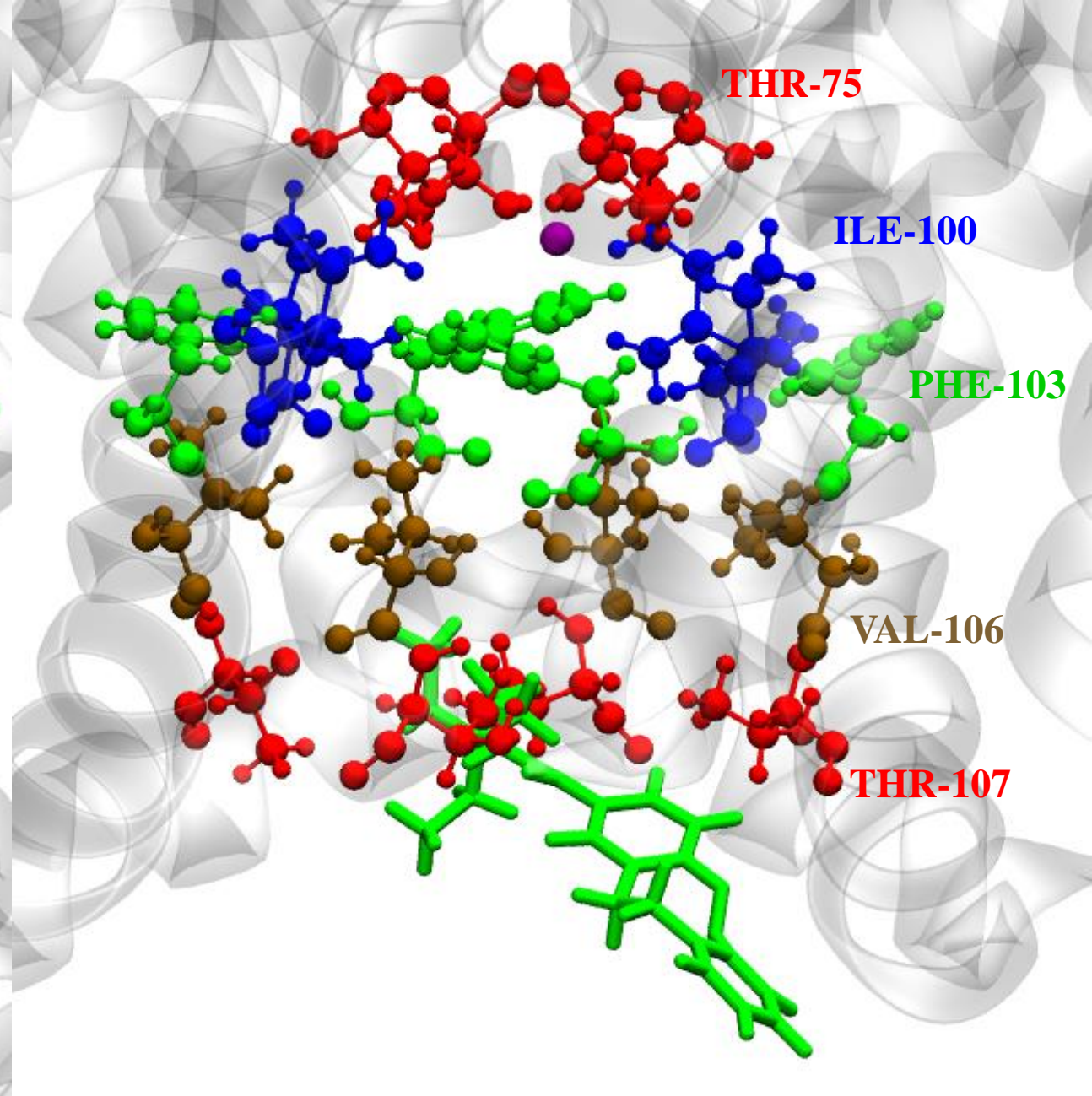


E1 (C0)
MM/PBSA Binding energy = -20.6 kcal/mol



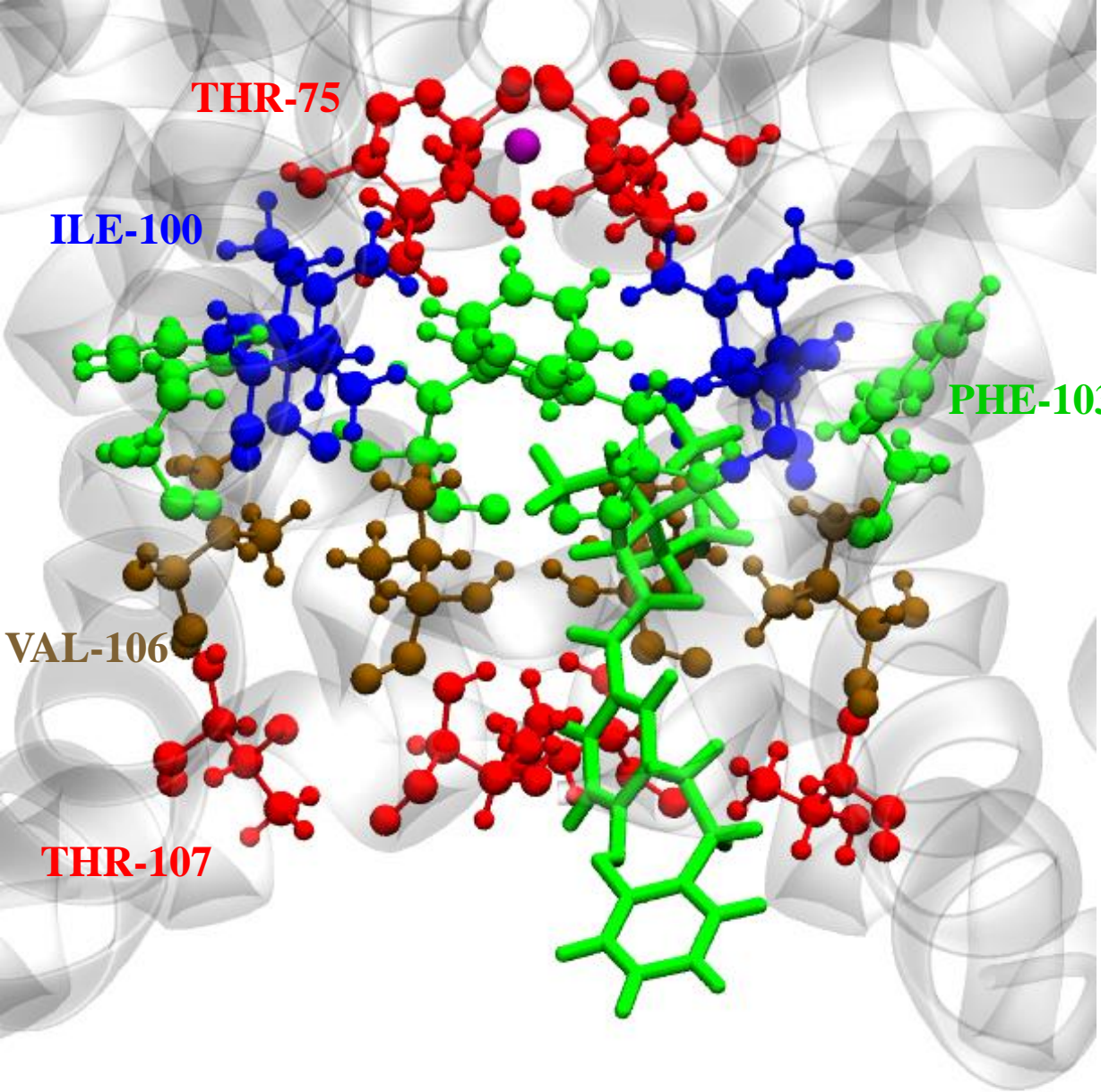
E1 (C1)

MM/PBSA Binding energy = -20.4 kcal/mol



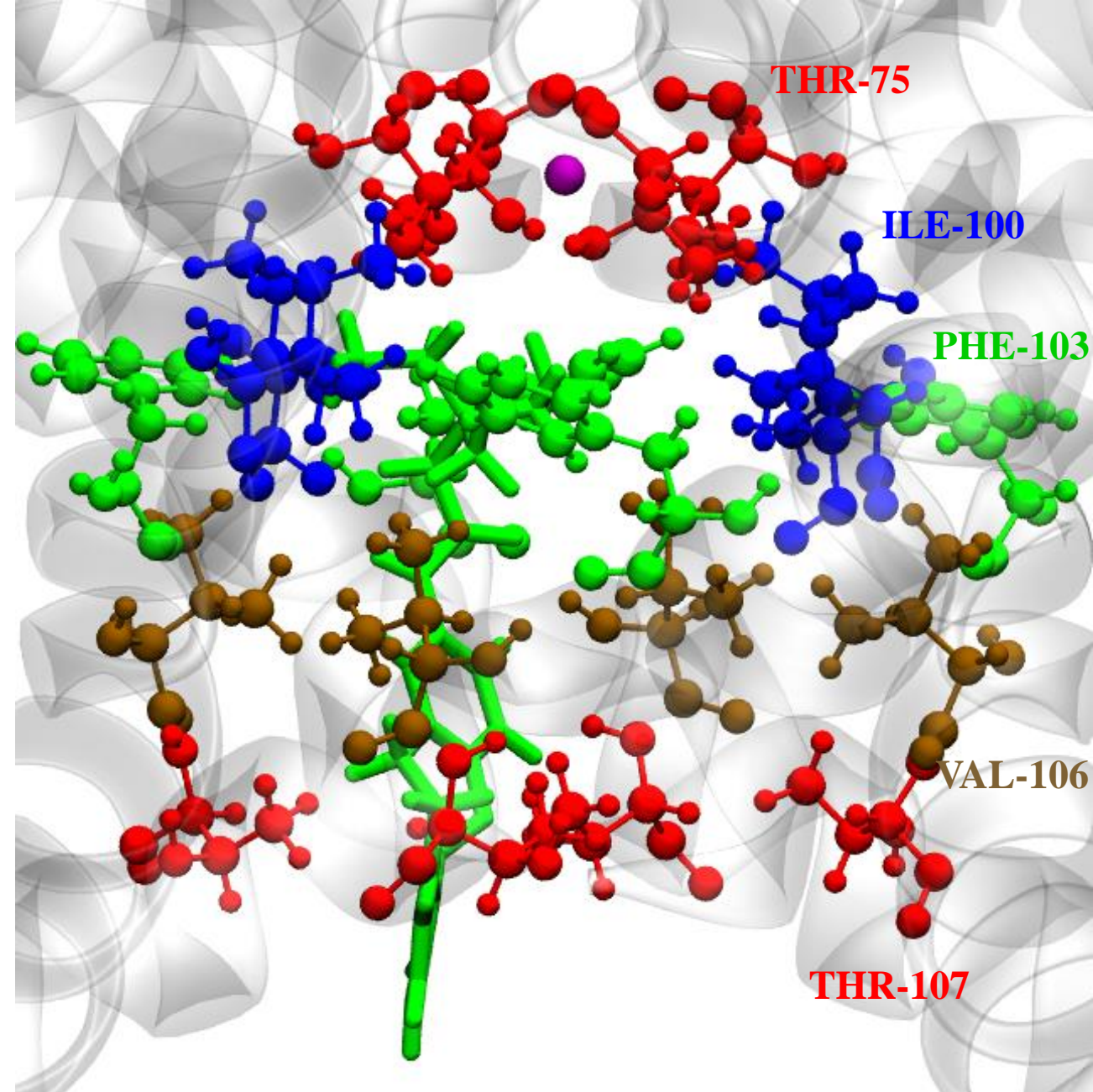
E1 (C2)

MM/PBSA Binding energy = -16.2 kcal/mol¹⁵



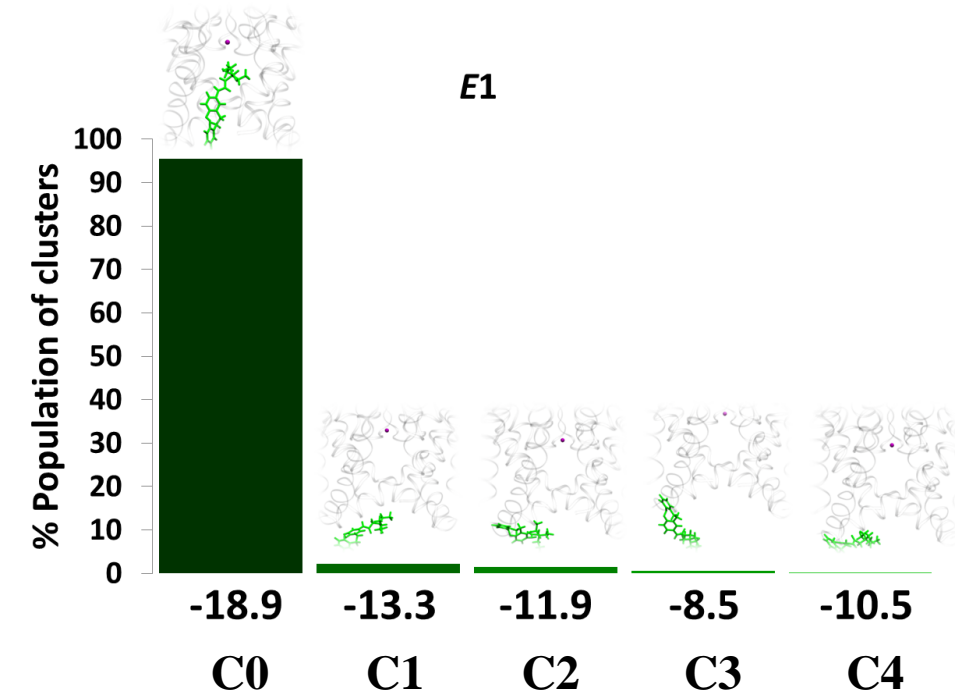
E1 (C3)

MM/PBSA Binding energy = -20.9 kcal/mol

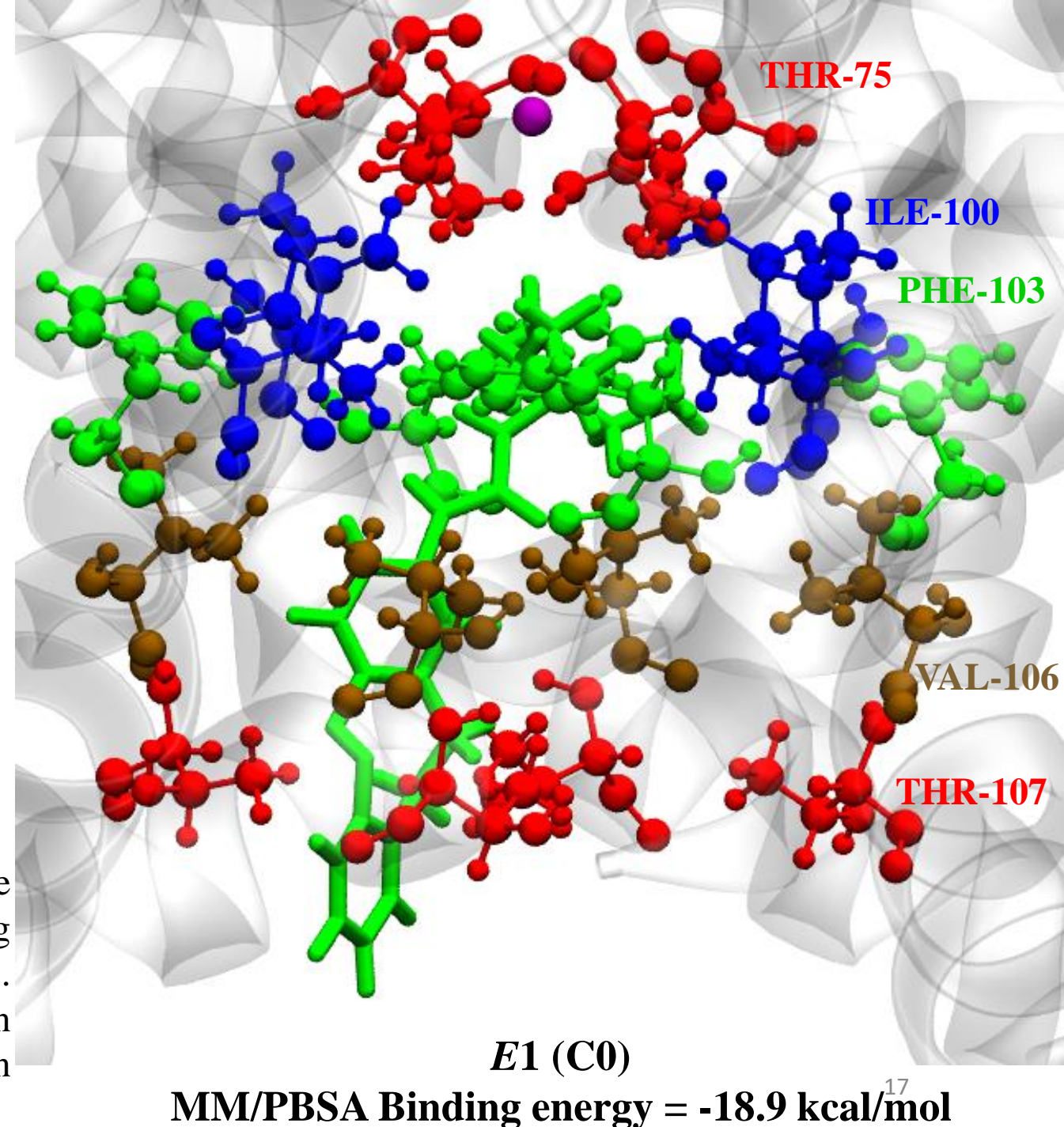


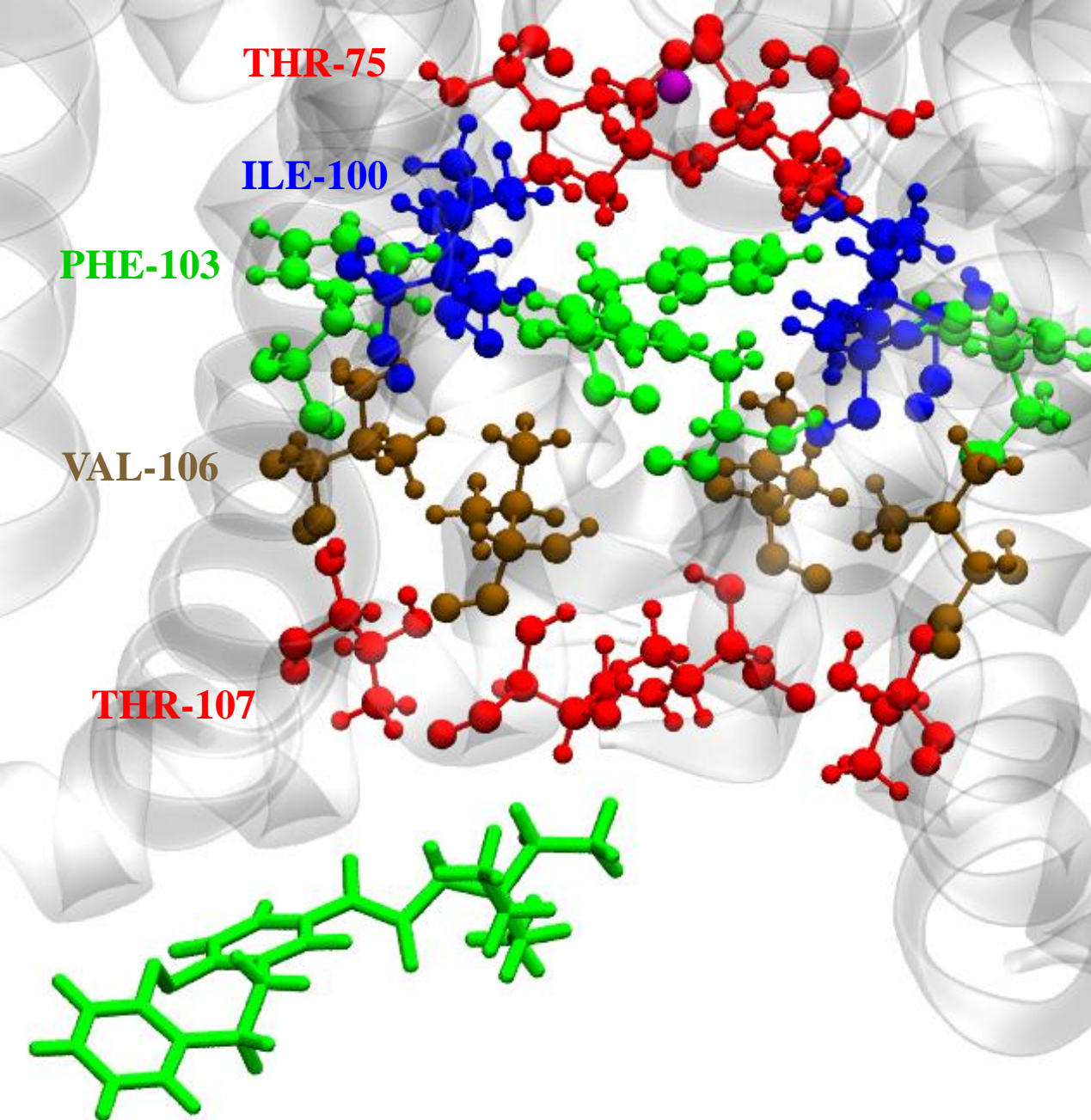
E1 (C4)

MM/PBSA Binding energy = -18.1 kcal/mol¹⁶



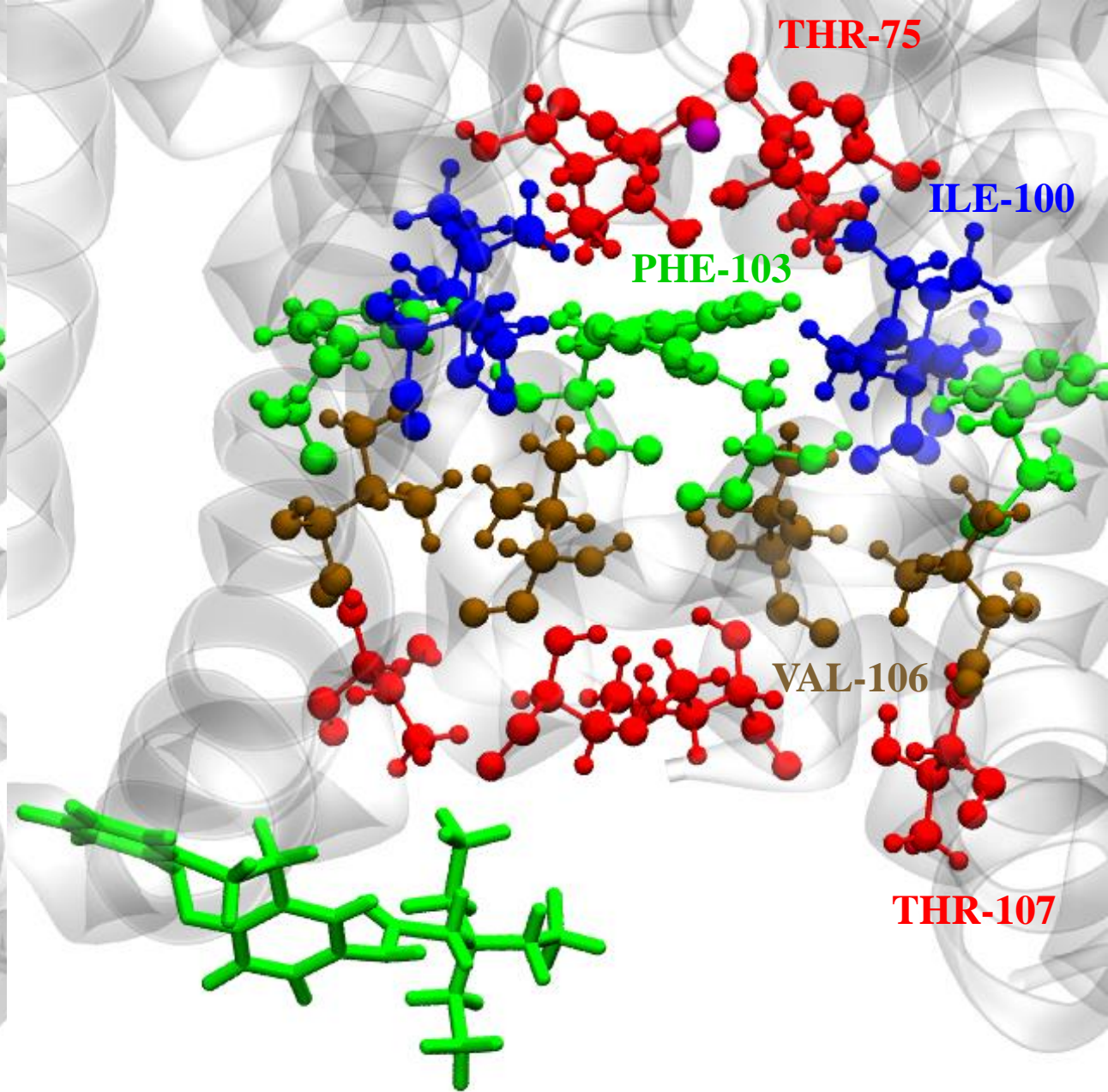
Cn (n = 0 – 1) indicate different clusters. The percentage population of clusters is given on the Y-axis. Corresponding binding energy (B.E.) is given in kcal/mol on the X-axis. Protein is represented in new cartoon model with K⁺-ion (purple ball), and E-LAB-TEA is shown in stick model (green colour).





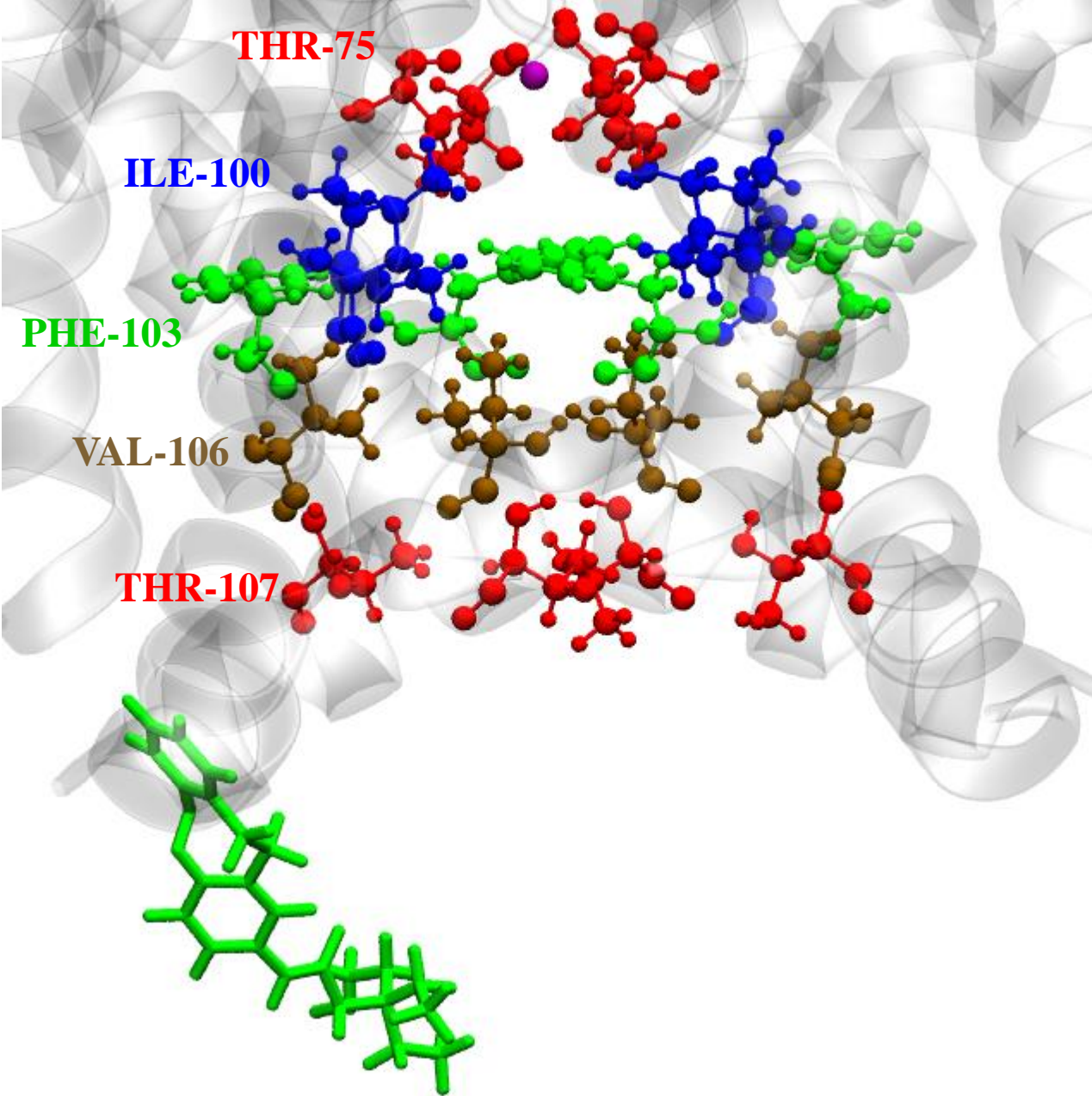
E1 (C1)

MM/PBSA Binding energy = -13.3 kcal/mol



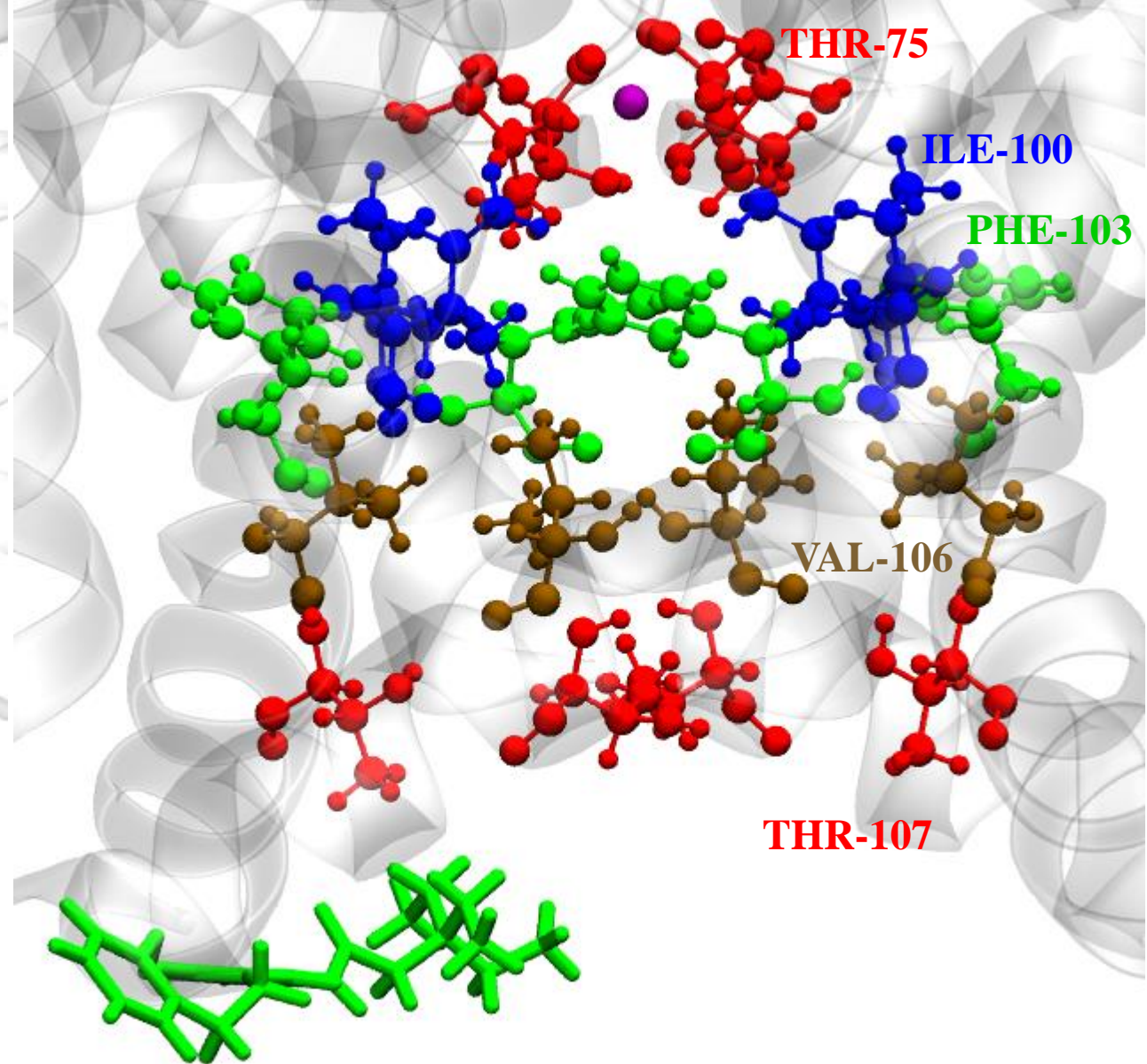
E1 (C2)

MM/PBSA Binding energy = -11.9 kcal/mol¹⁸



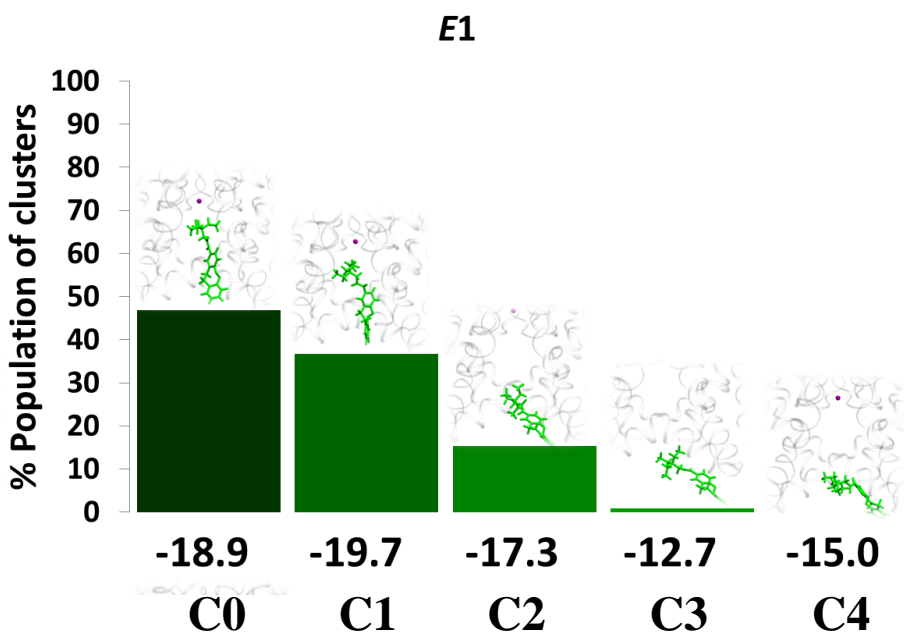
E1 (C3)

MM/PBSA Binding energy = -8.5 kcal/mol

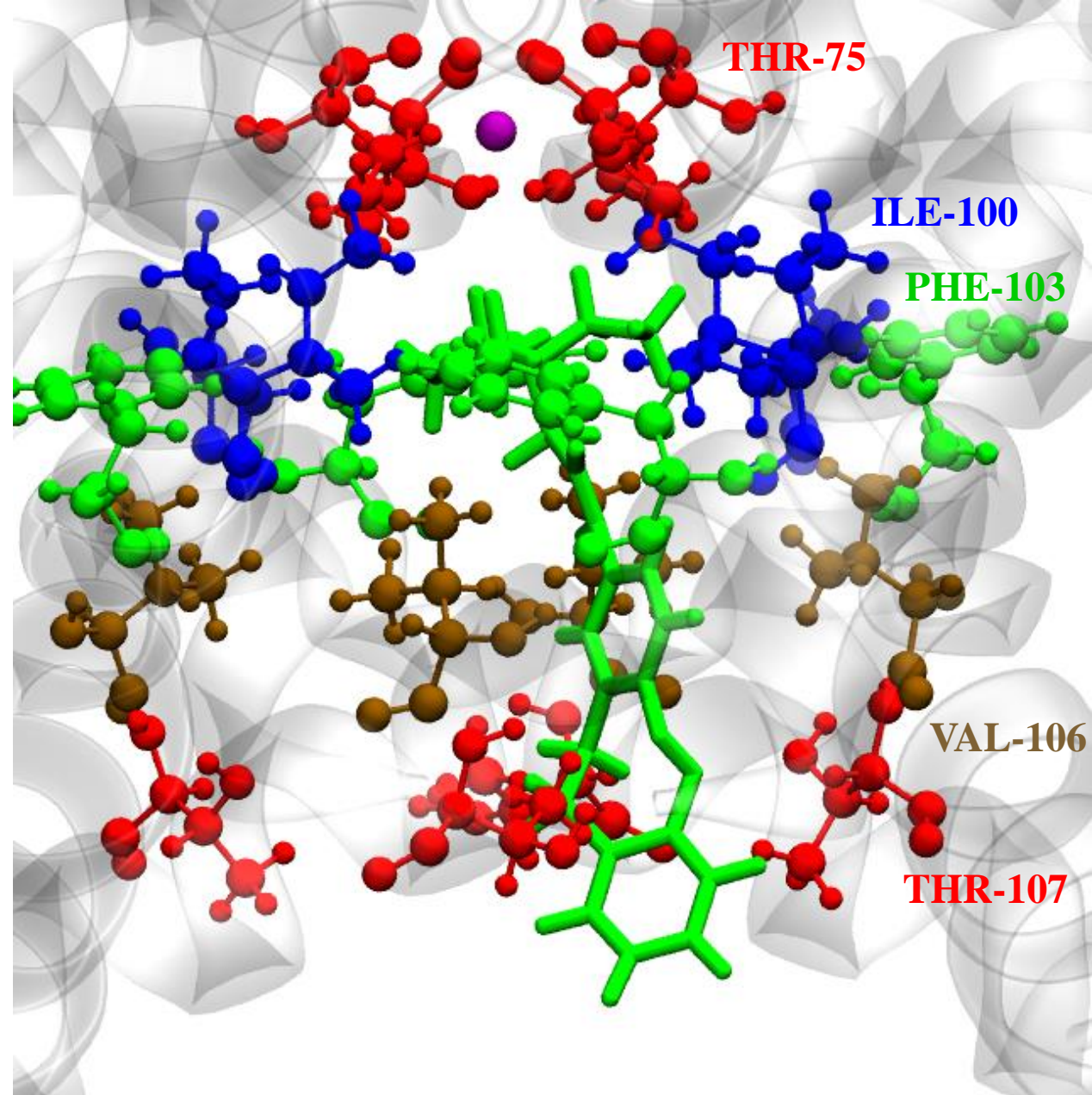


E1 (C4)

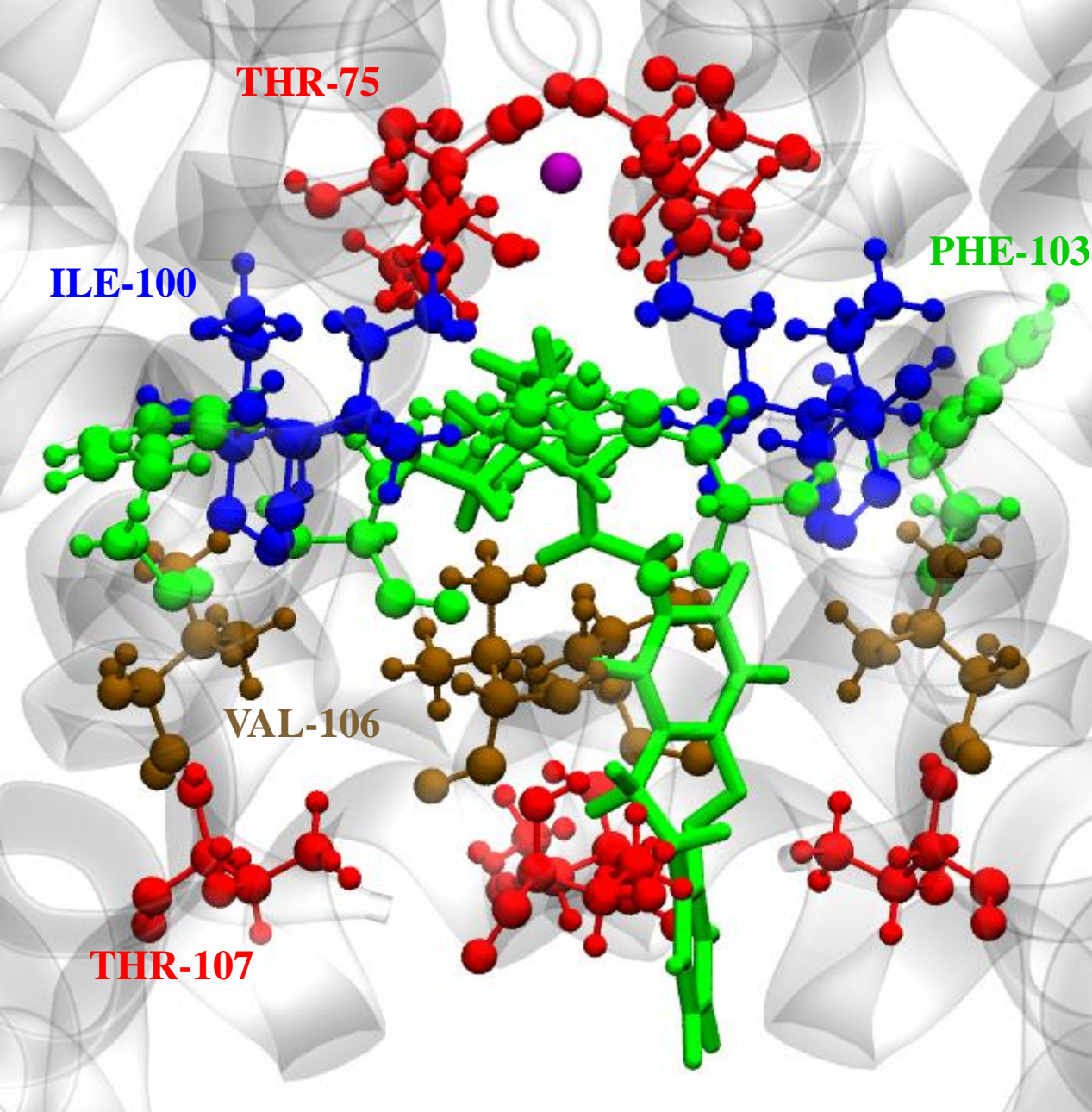
MM/PBSA Binding energy = -10.5 kcal/mol¹⁹



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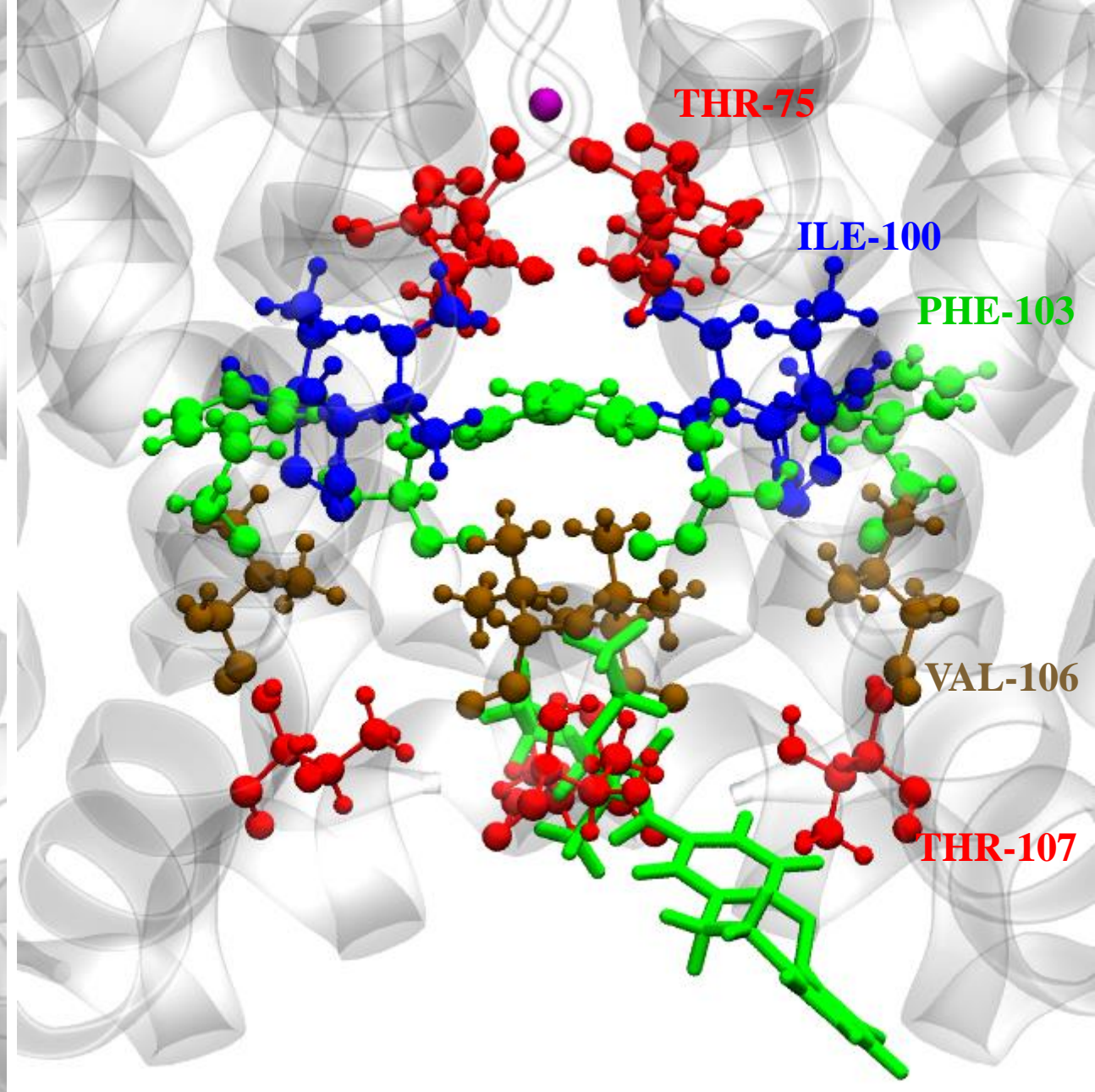


E1 (C0)
MM/PBSA Binding energy = -18.9 kcal/mol²⁰



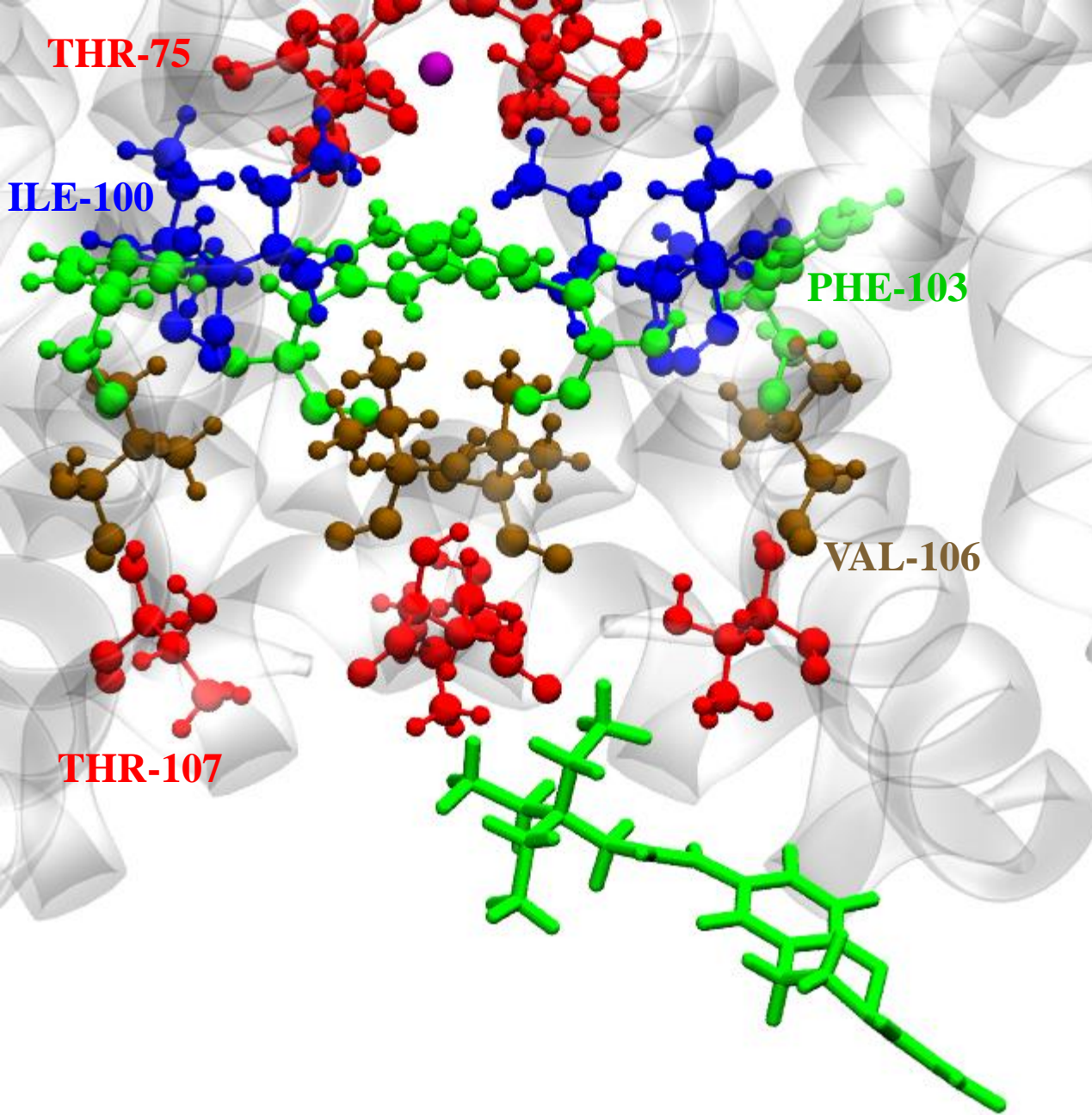
E1 (C1)

MM/PBSA Binding energy = -19.7 kcal/mol



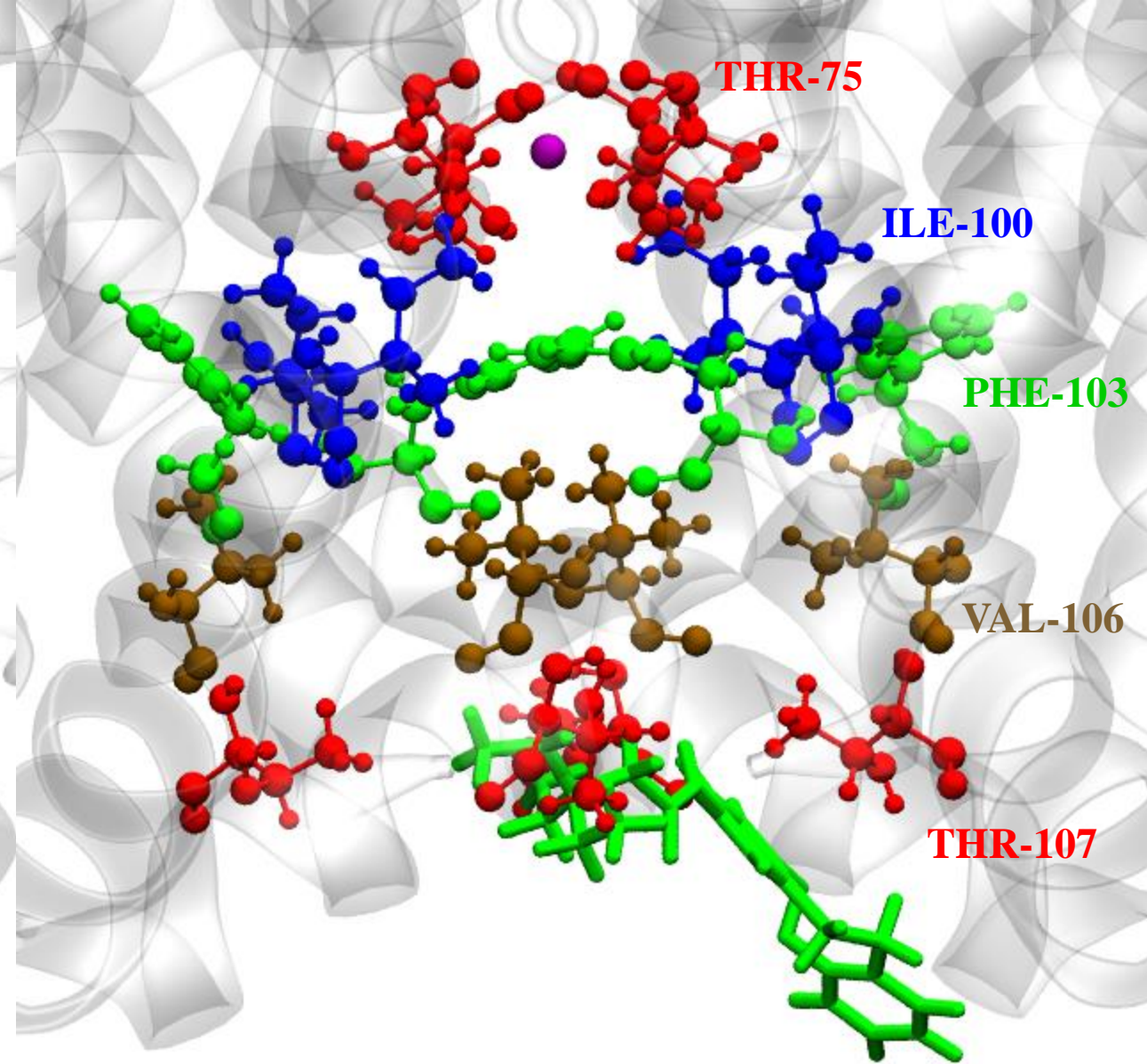
E1 (C2)

MM/PBSA Binding energy = -17.3 kcal/mol²¹



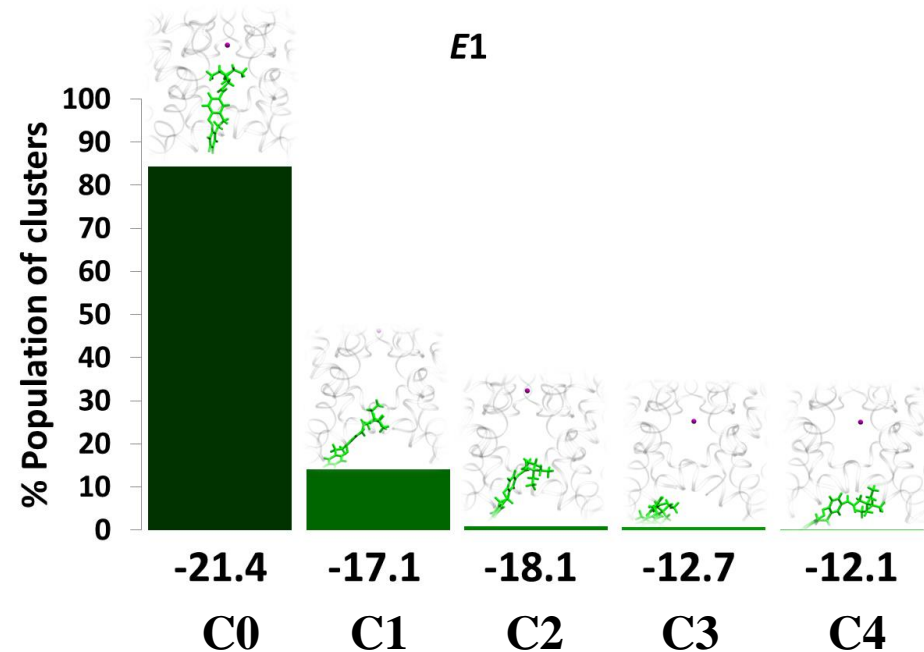
E1 (C3)

MM/PBSA Binding energy = -12.7 kcal/mol

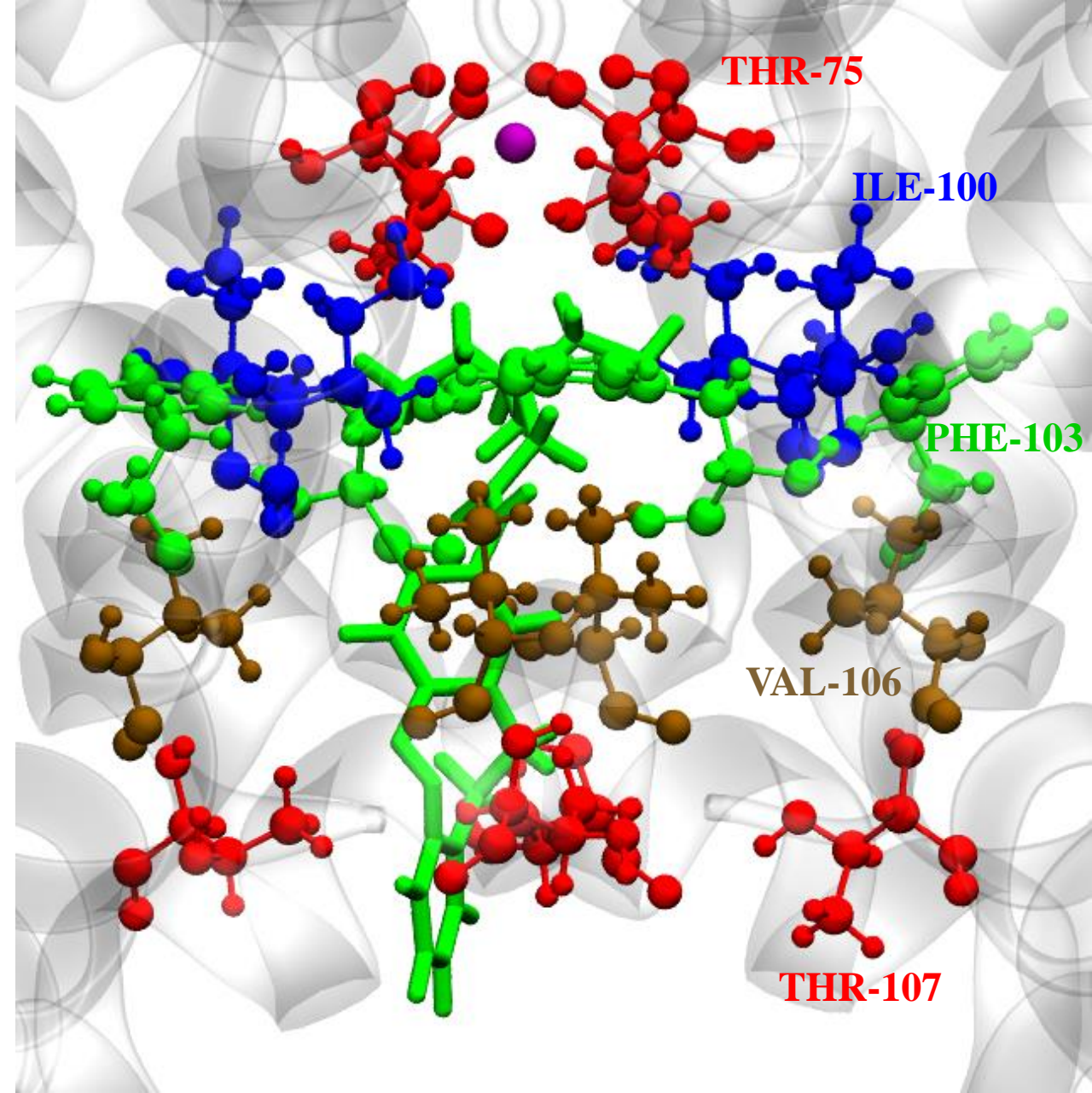


E1 (C4)

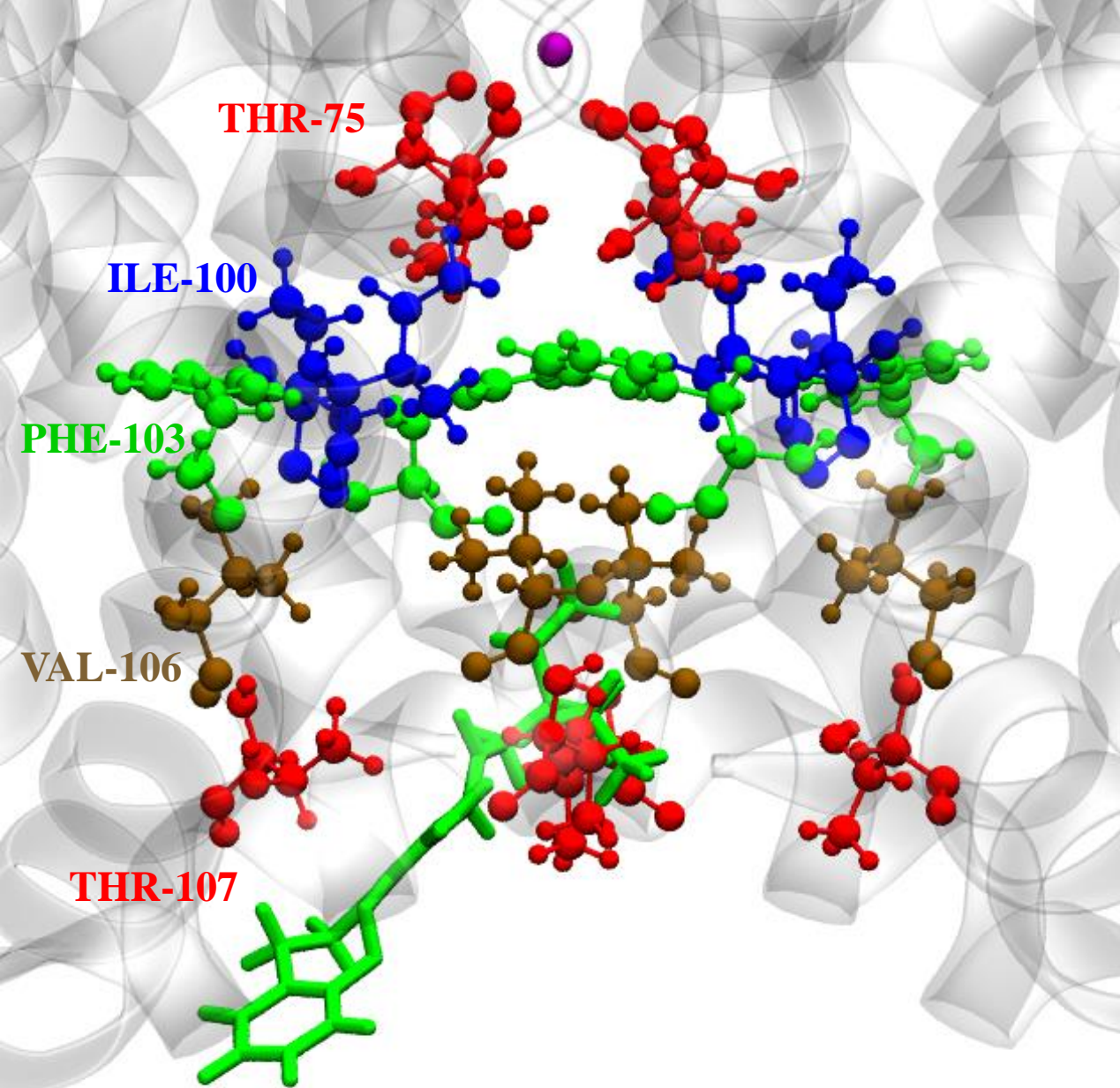
MM/PBSA Binding energy = -15.0 kcal/mol²²



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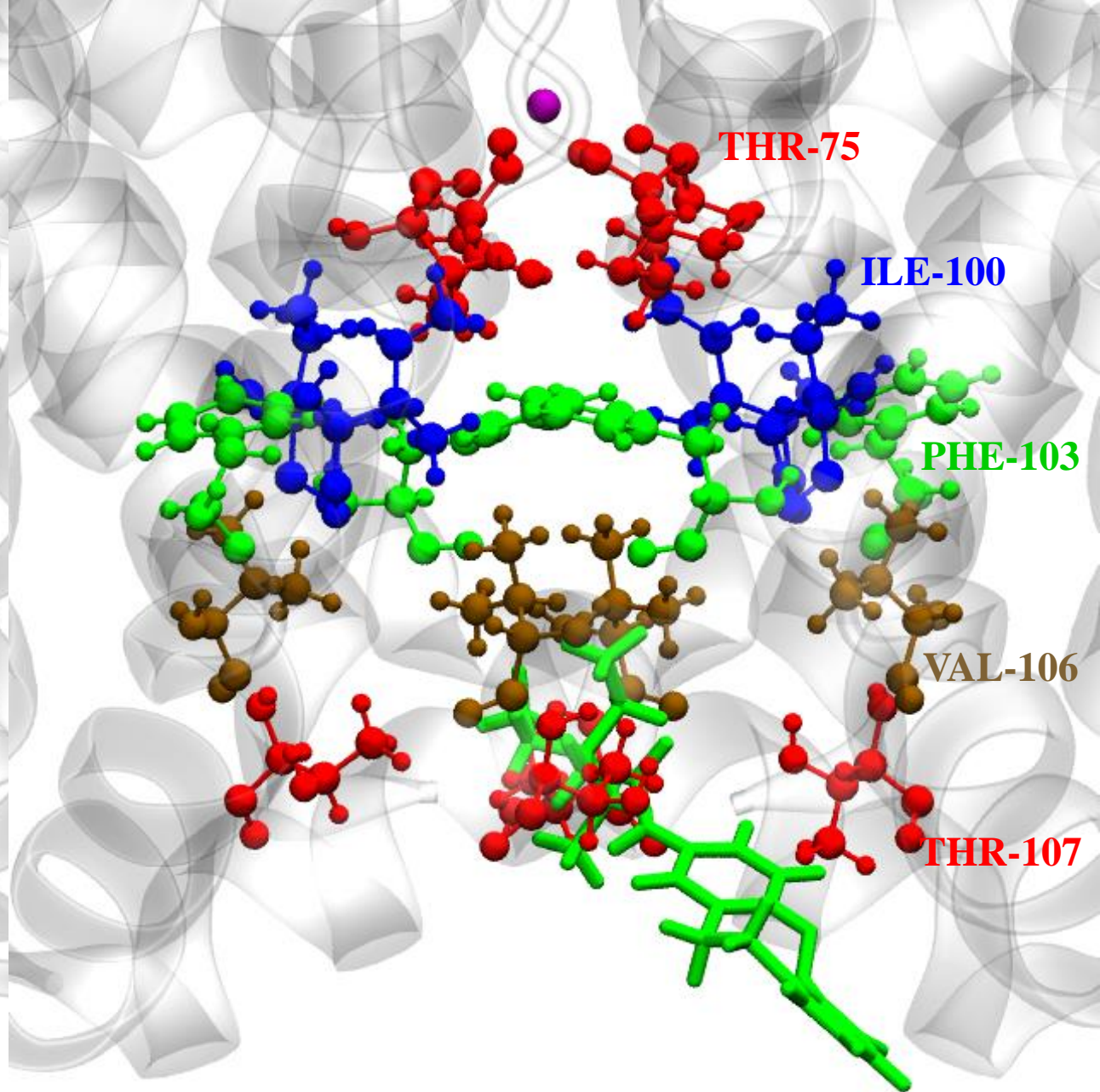


***E*1 (C0)**
MM/PBSA Binding energy = -21.4 kcal/mol²³



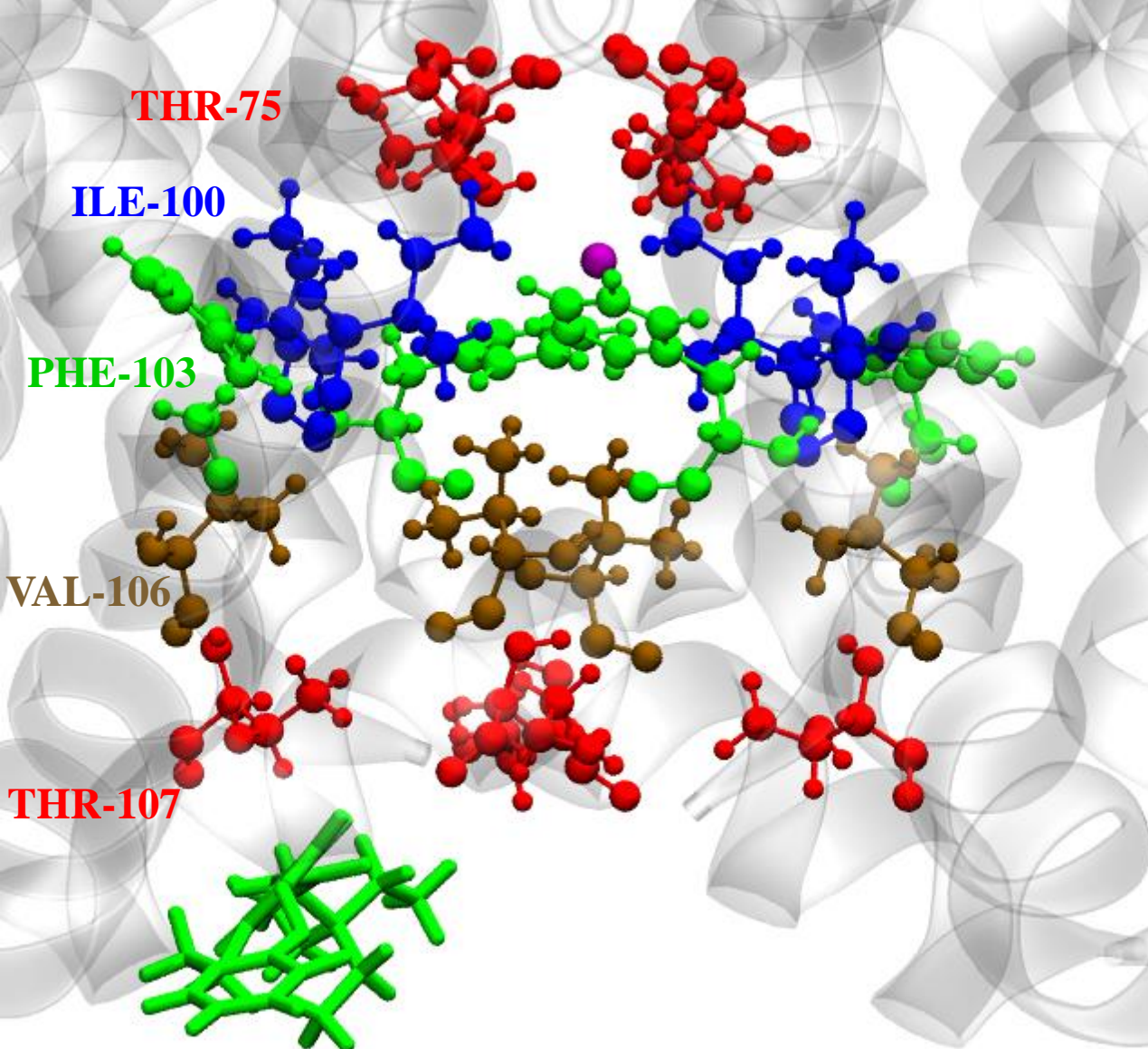
E1 (C1)

MM/PBSA Binding energy = -17.1 kcal/mol



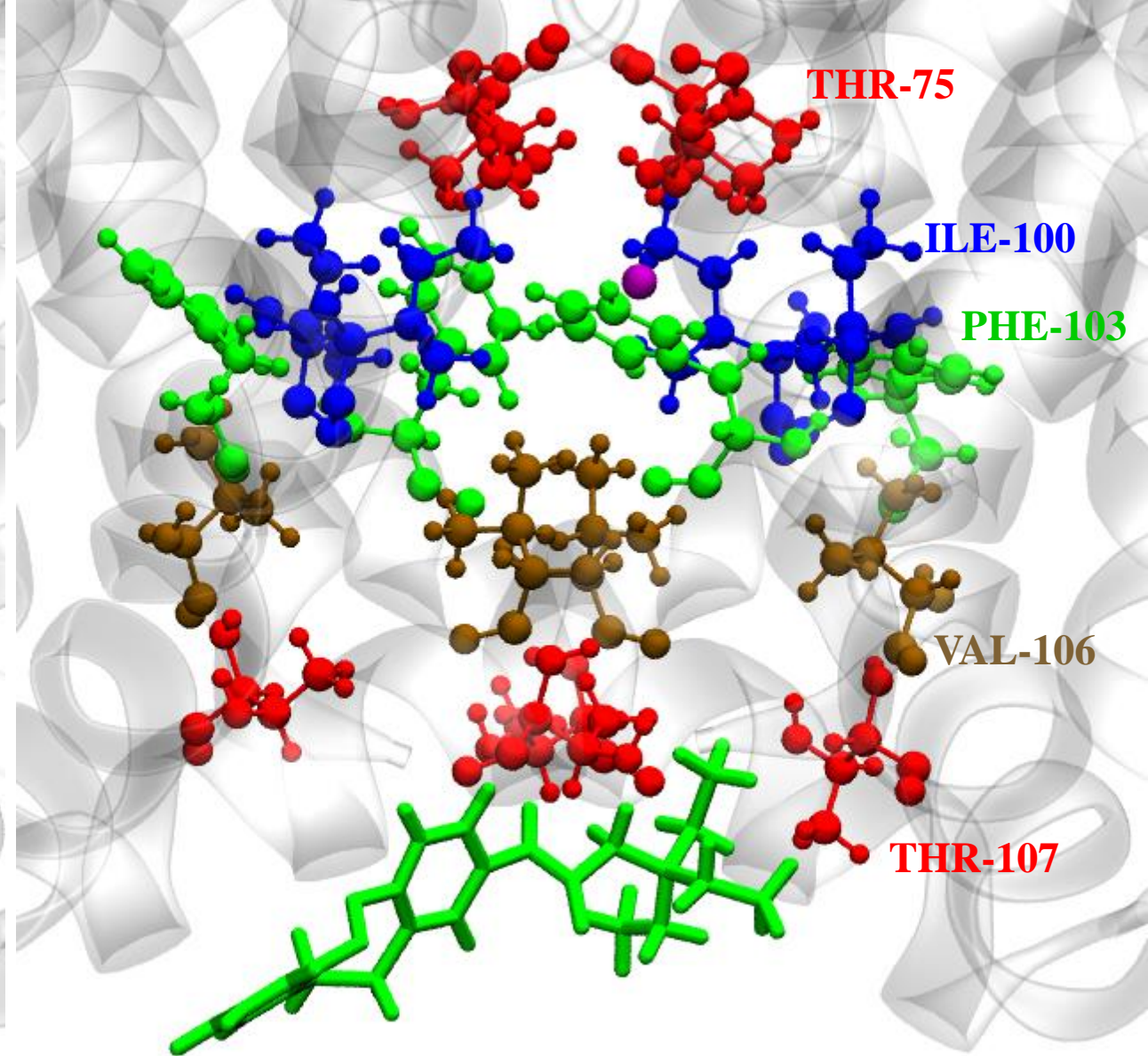
E1 (C2)

MM/PBSA Binding energy = -18.1 kcal/mol²⁴



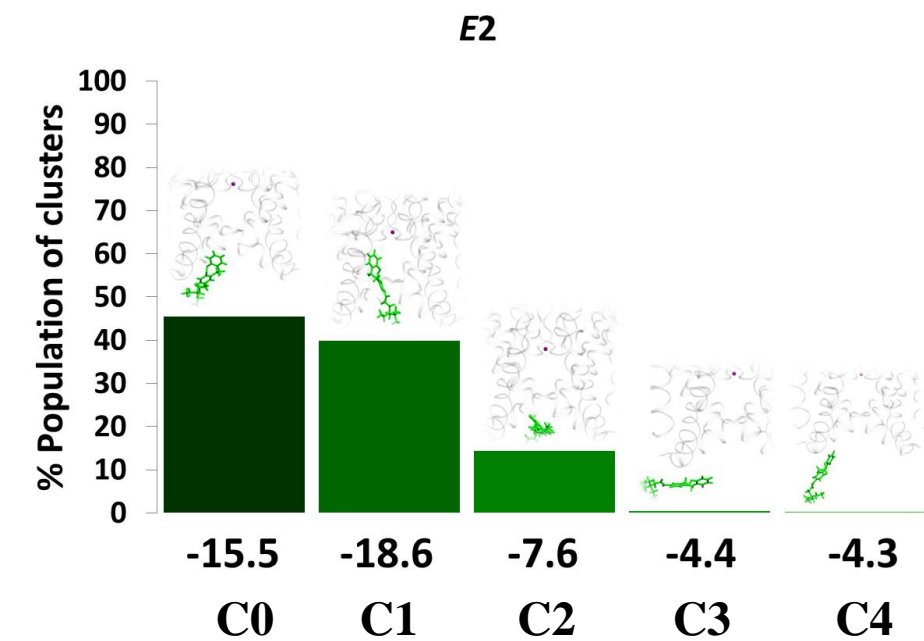
E1 (C3)

MM/PBSA Binding energy = -12.7 kcal/mol

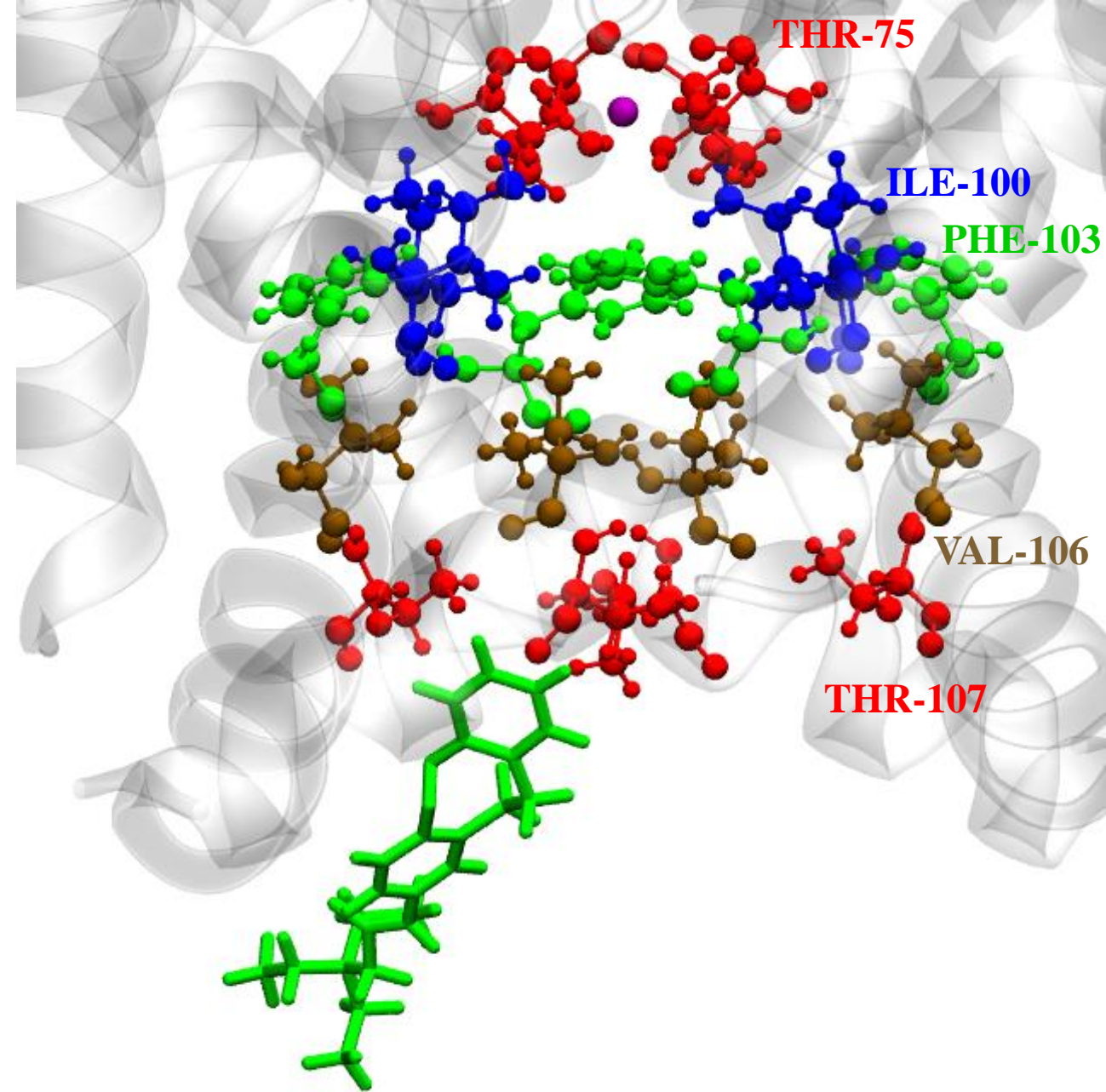


E1 (C4)

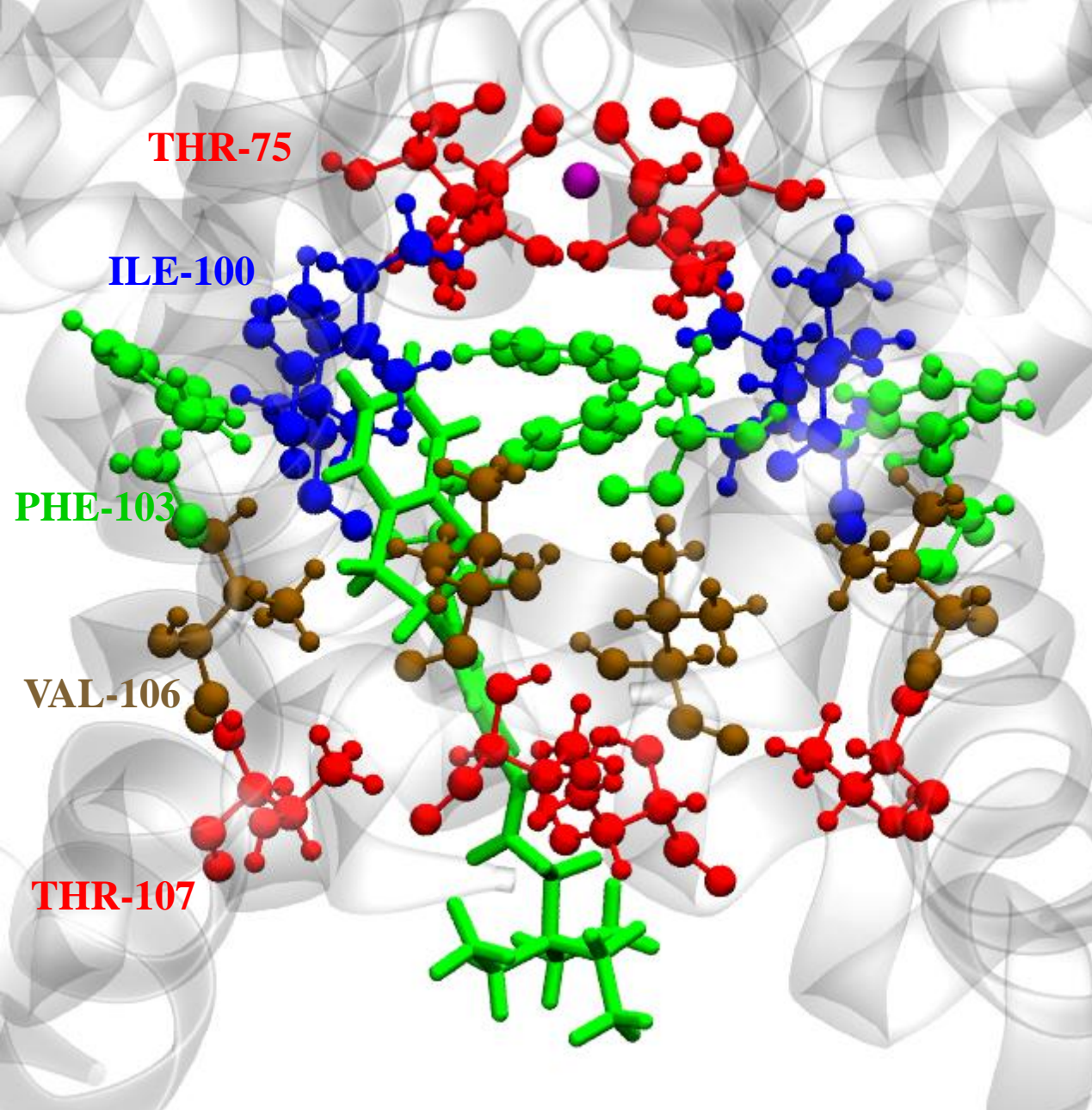
MM/PBSA Binding energy = -12.1 kcal/mol²⁵



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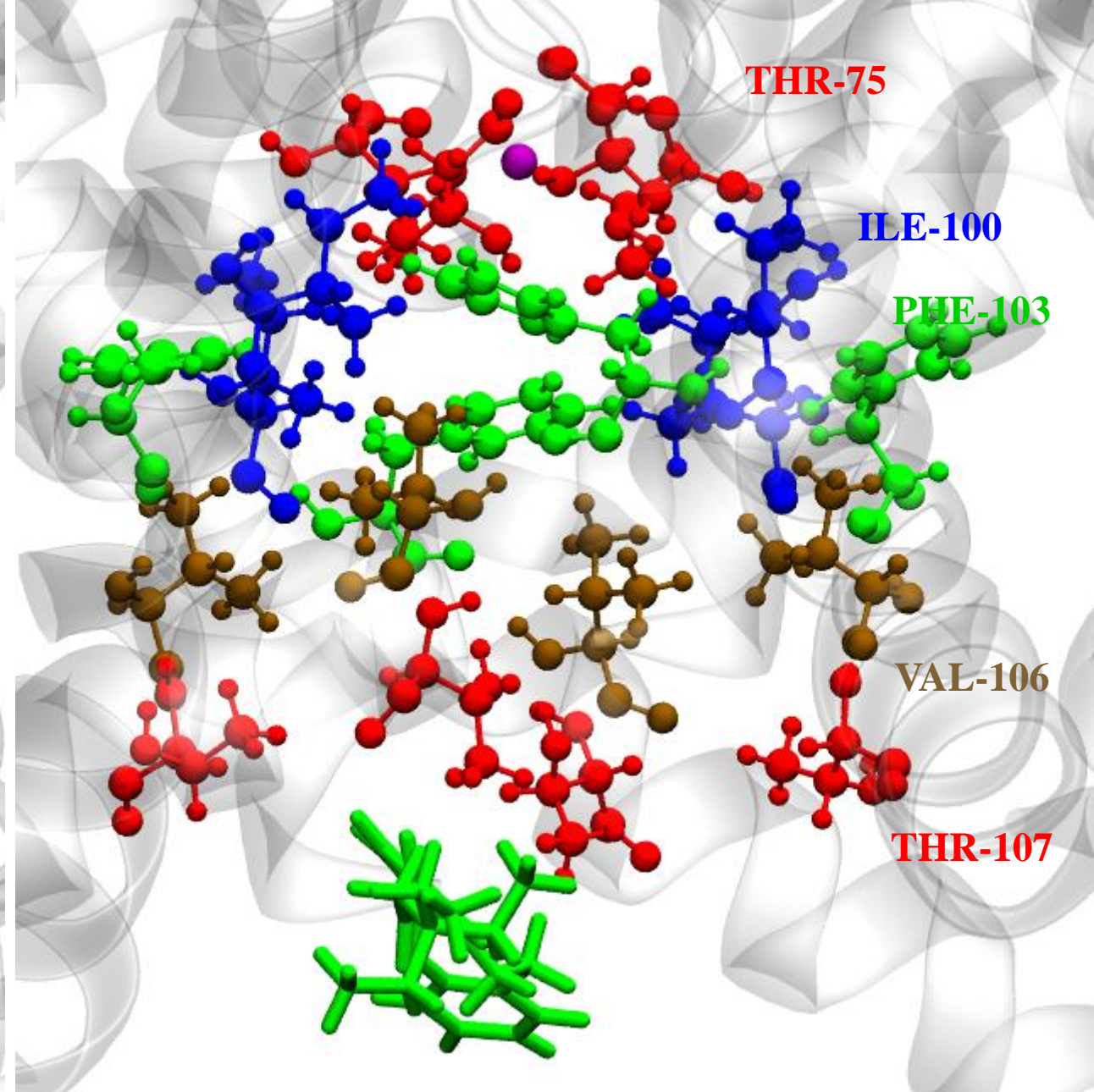


***E2* (C0)**
MM/PBSA Binding energy = -15.5 kcal/mol²⁶



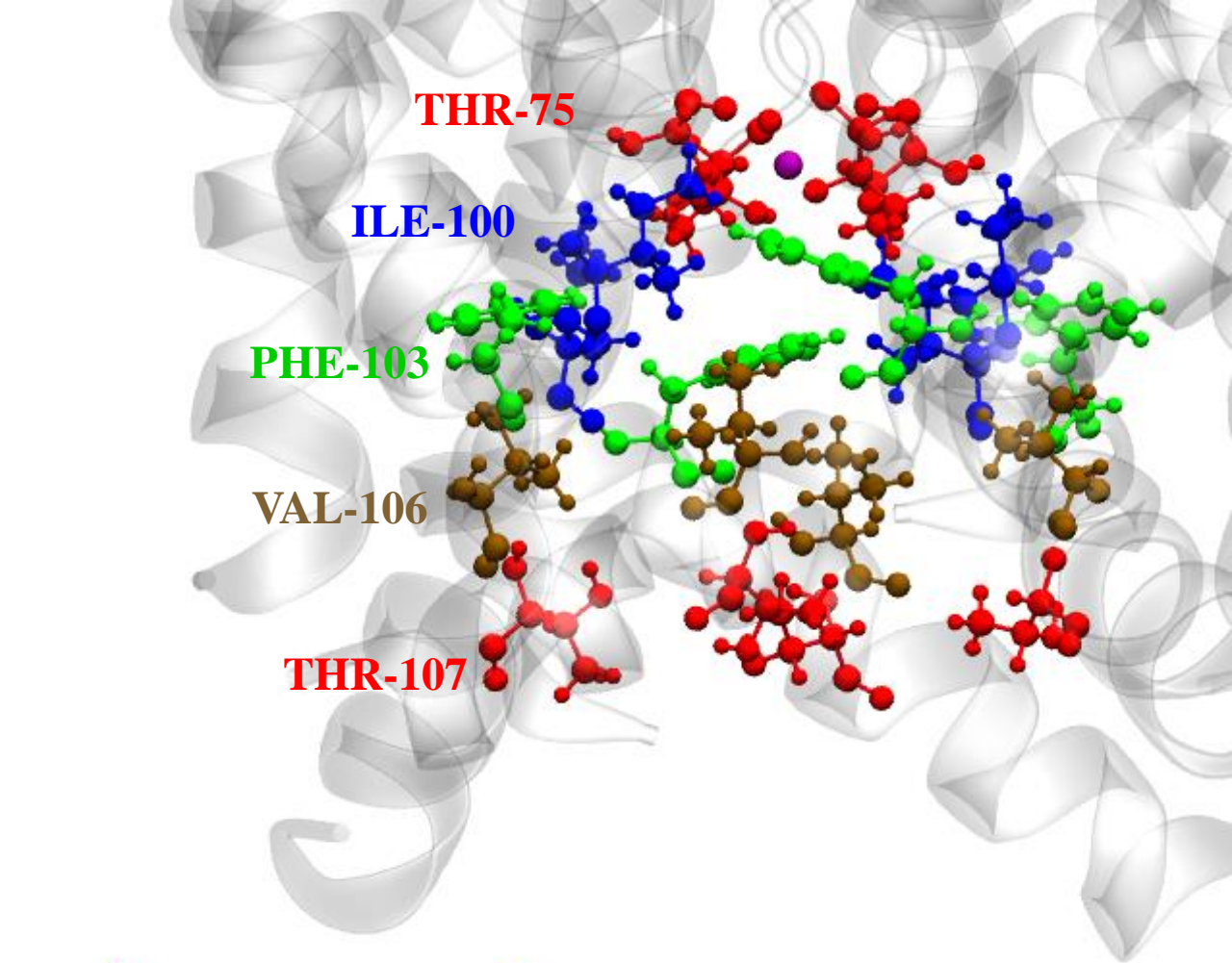
E2 (C1)

MM/PBSA Binding energy = -18.6 kcal/mol



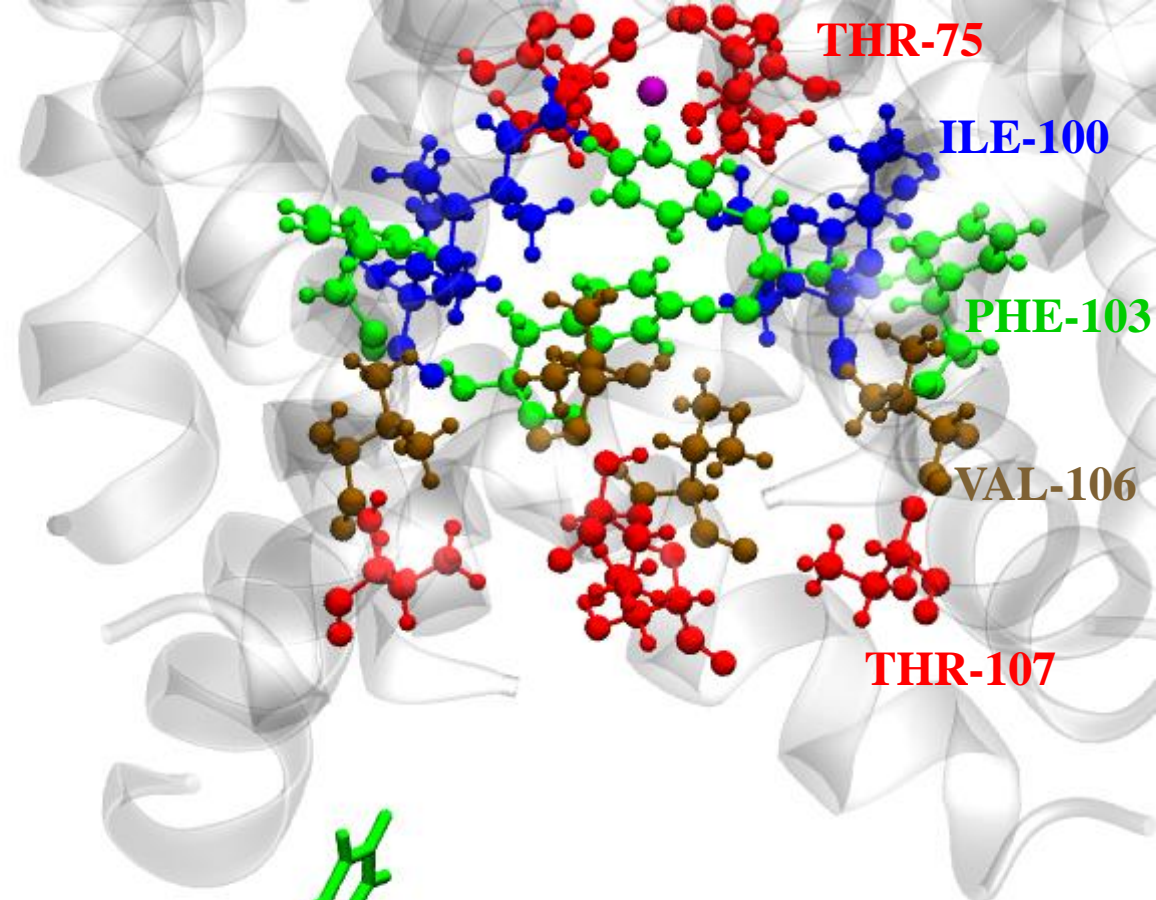
E2 (C2)

MM/PBSA Binding energy = -7.6 kcal/mol²⁷



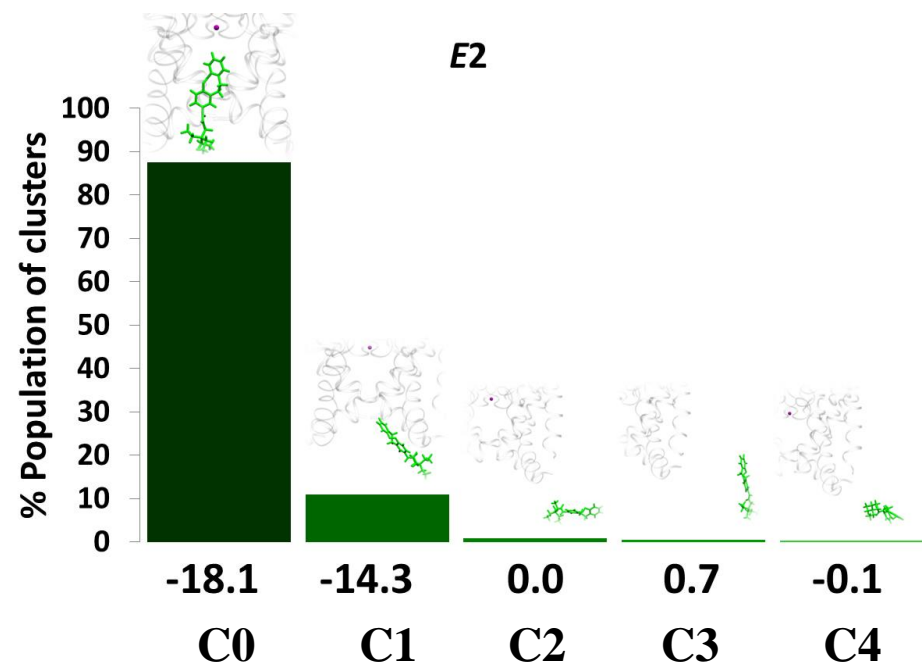
E2 (C3)

MM/PBSA Binding energy = -4.4 kcal/mol

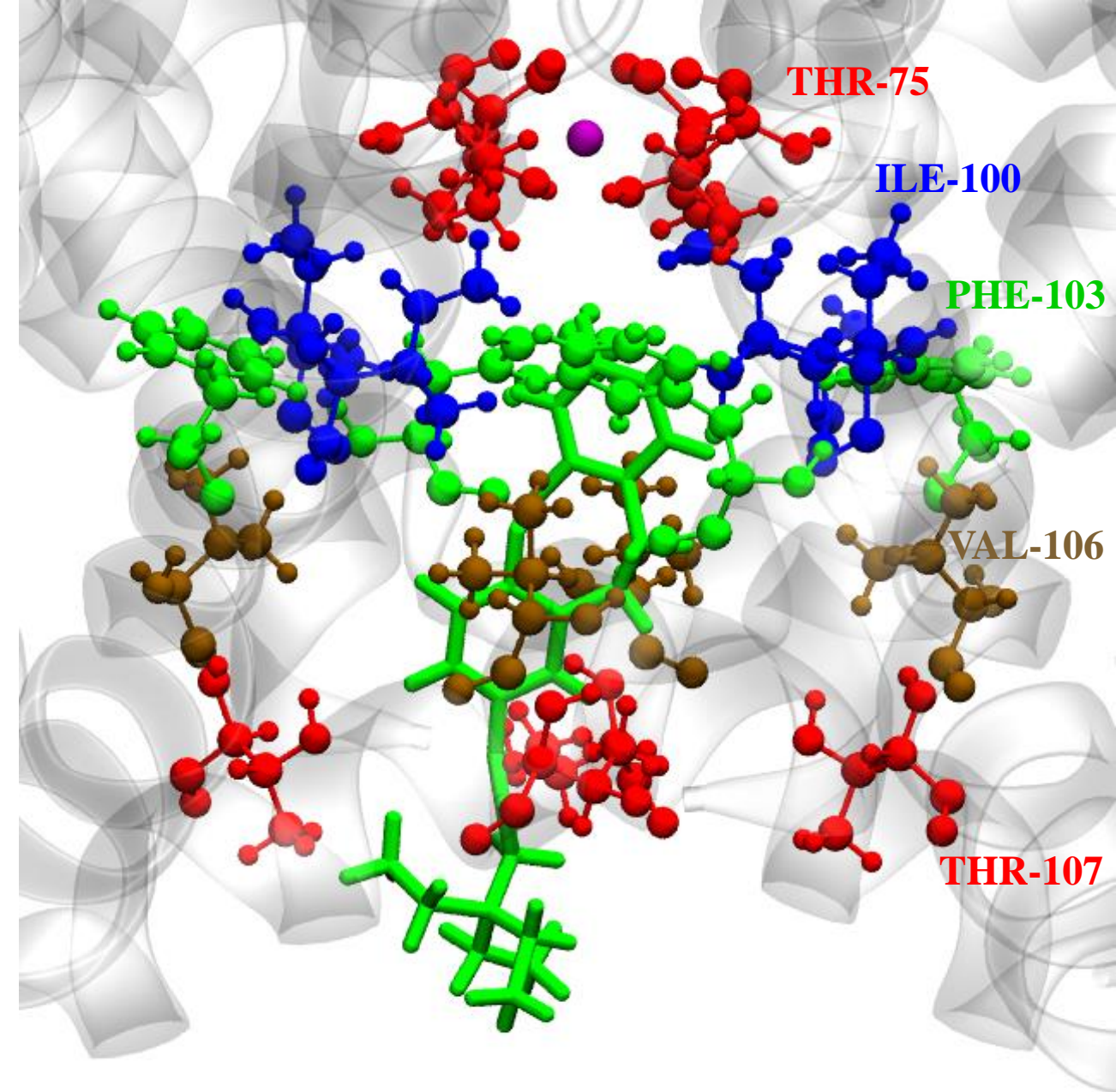


E2 (C4)

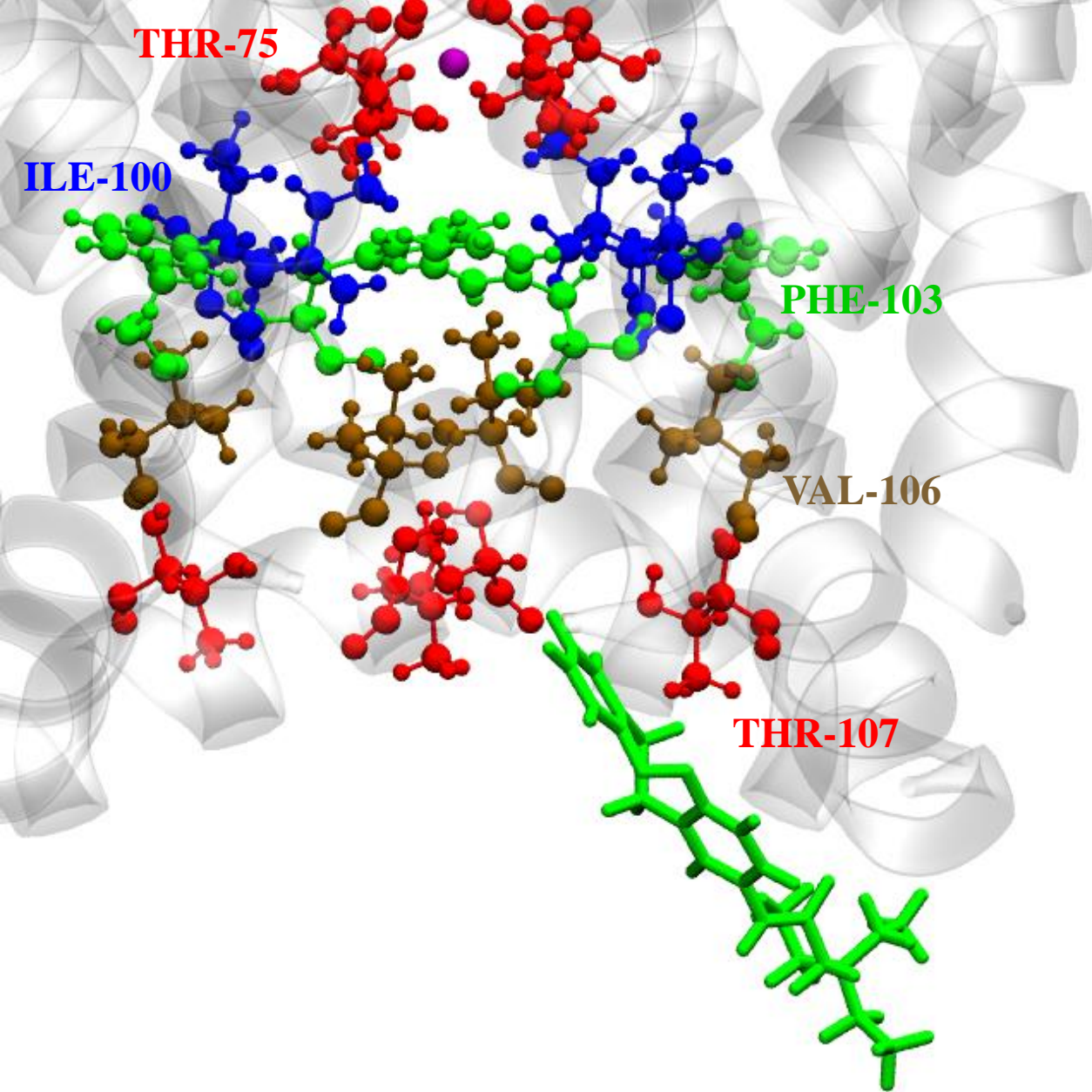
MM/PBSA Binding energy = -4.3 kcal/mol²⁸



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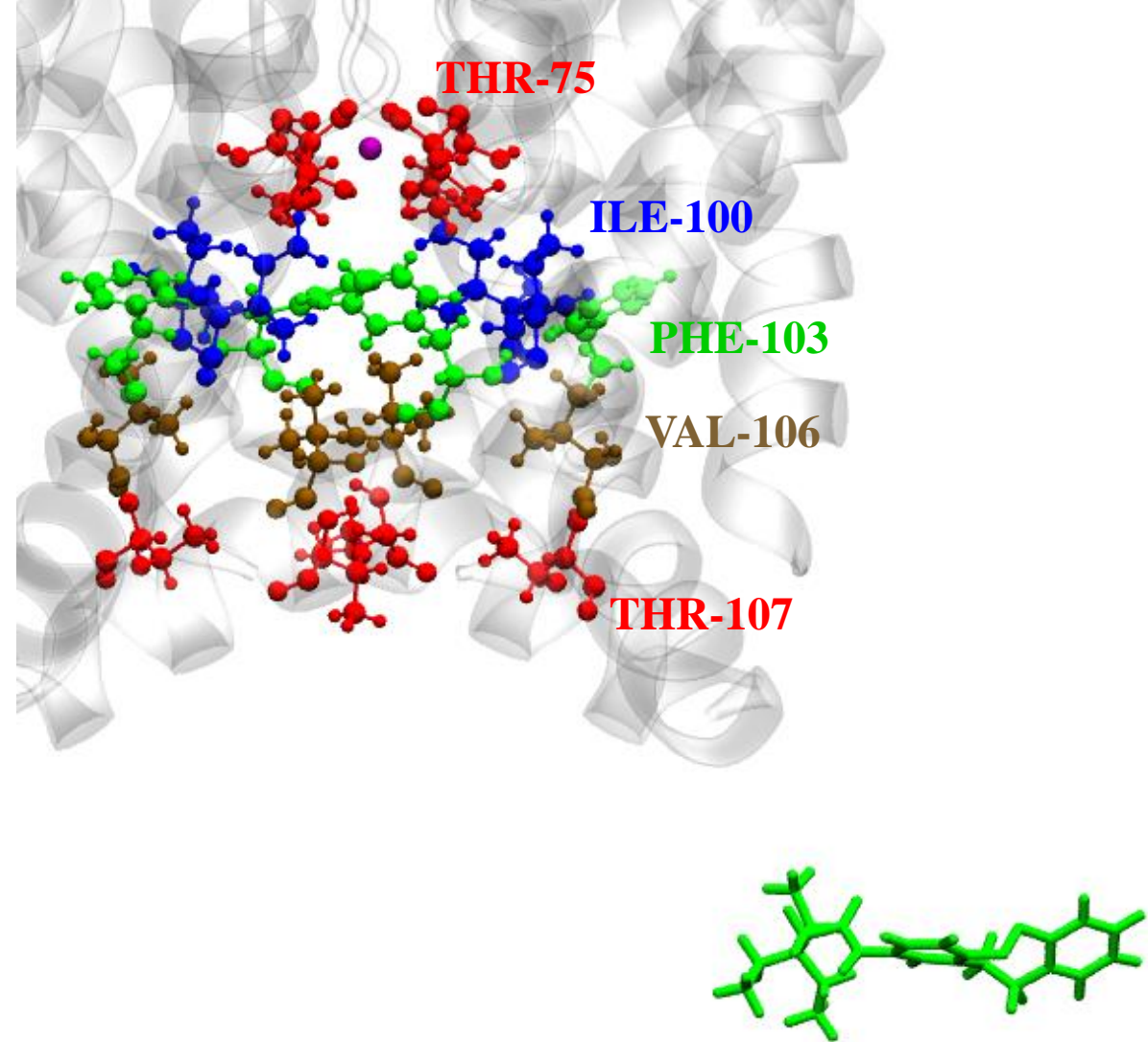


***E2* (C0)**
MM/PBSA Binding energy = -18.1 kcal/mol²⁹



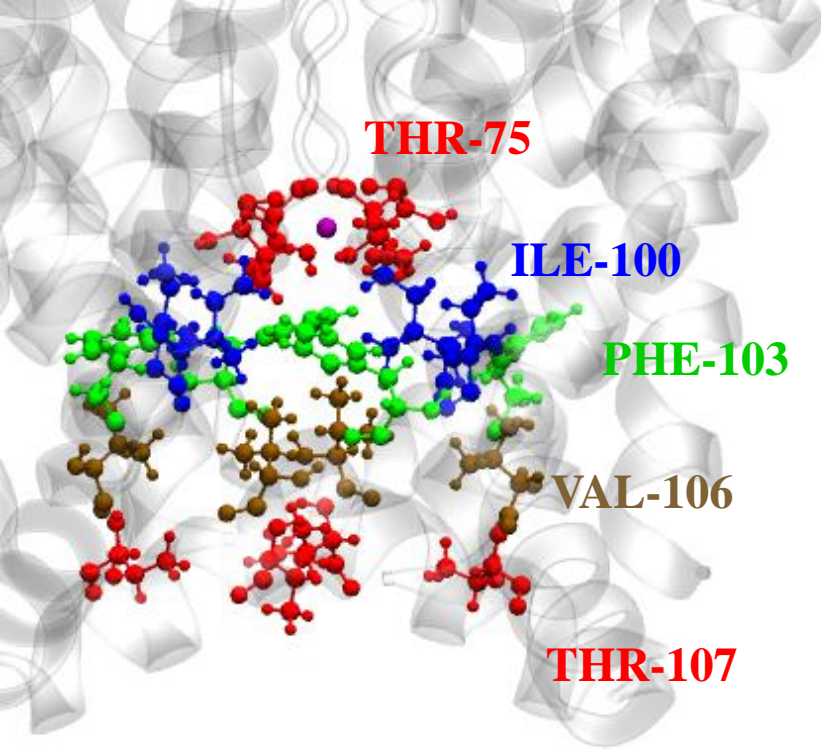
E2 (C1)

MM/PBSA Binding energy = -14.3 kcal/mol



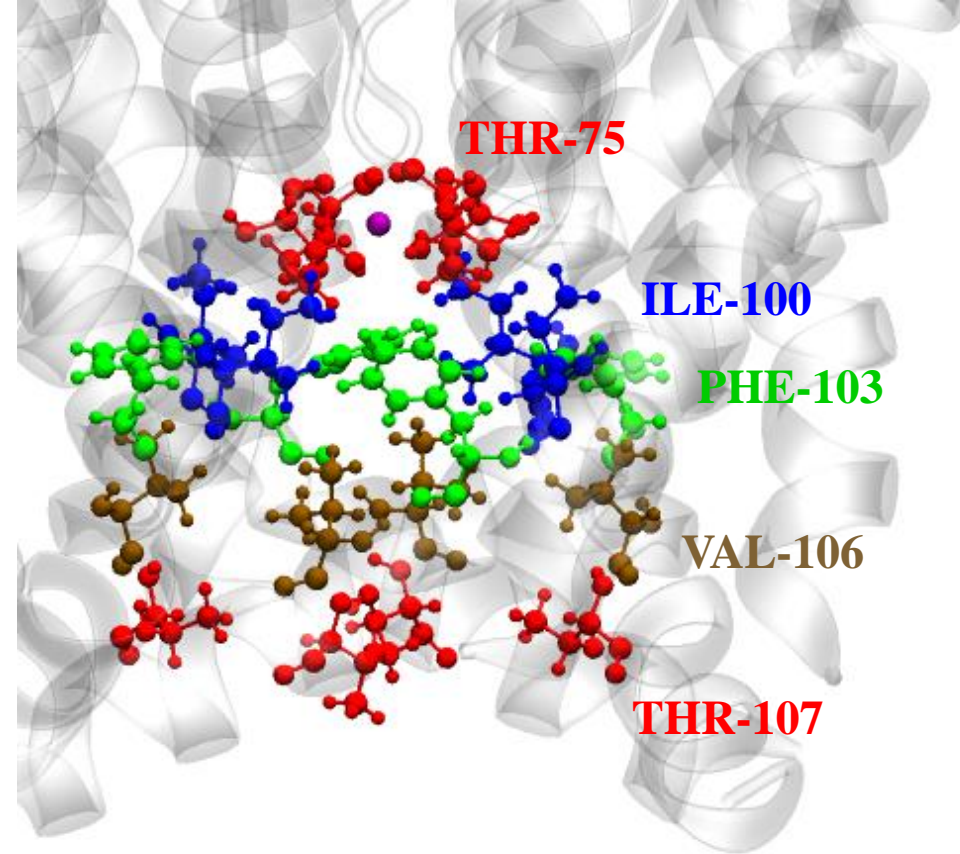
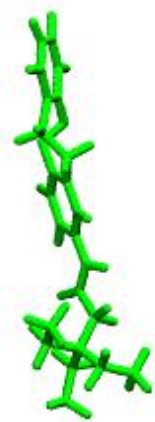
E2 (C2)

MM/PBSA Binding energy = 0.0 kcal/mol



E2 (C3)

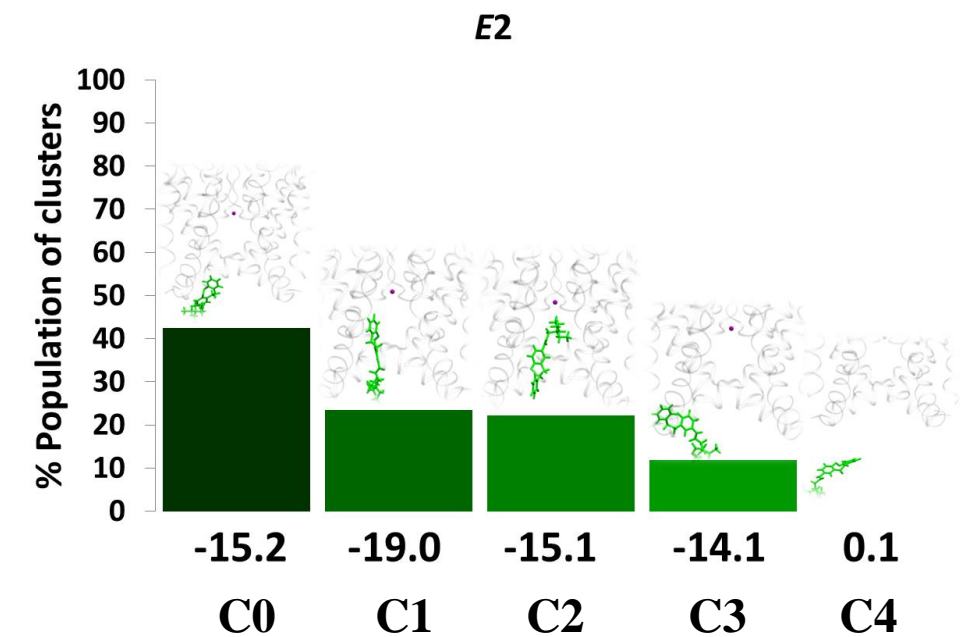
MM/PBSA Binding energy = 0.7 kcal/mol



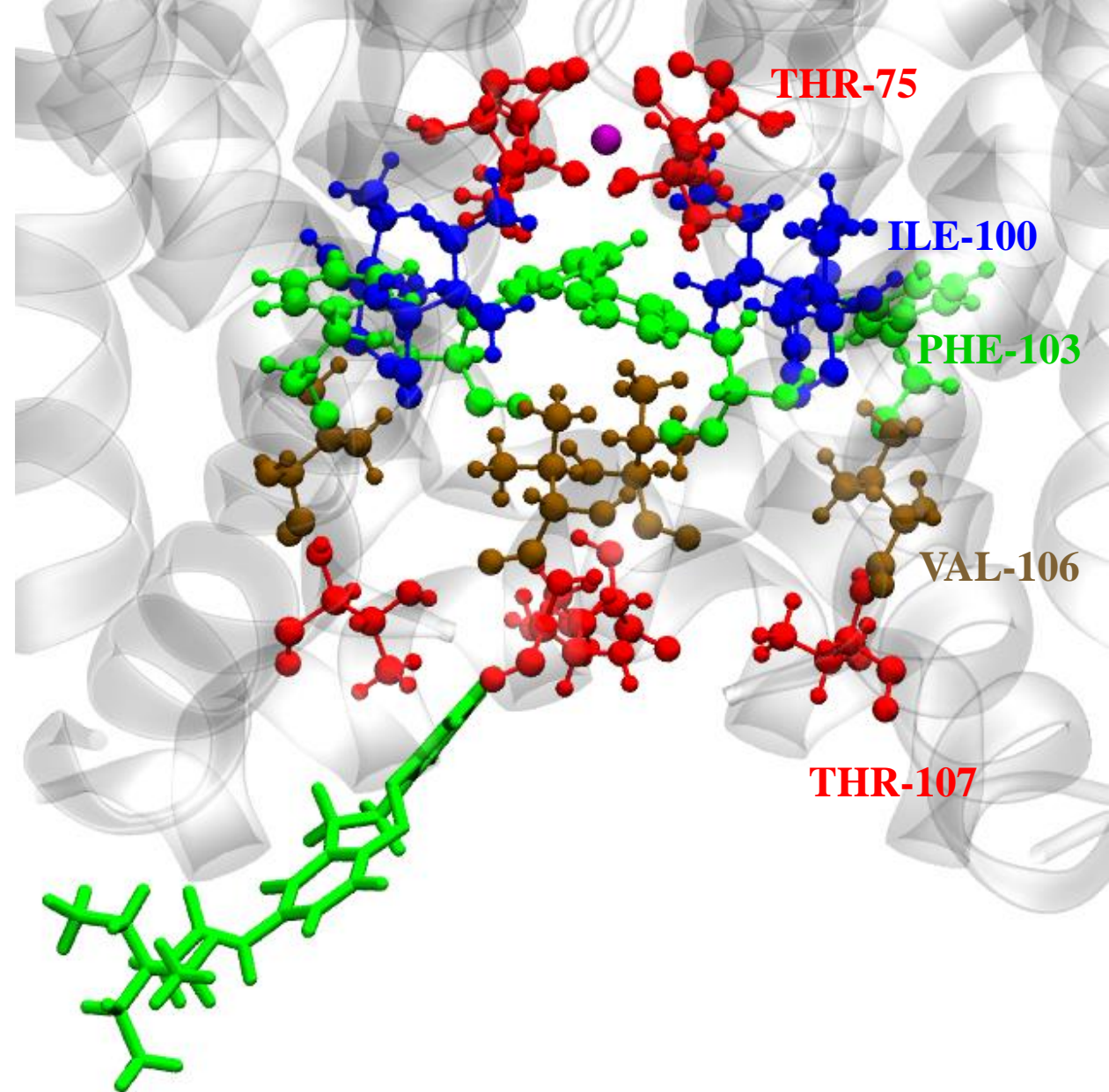
E2 (C4)

MM/PBSA Binding energy = -0.1 kcal/mol³¹

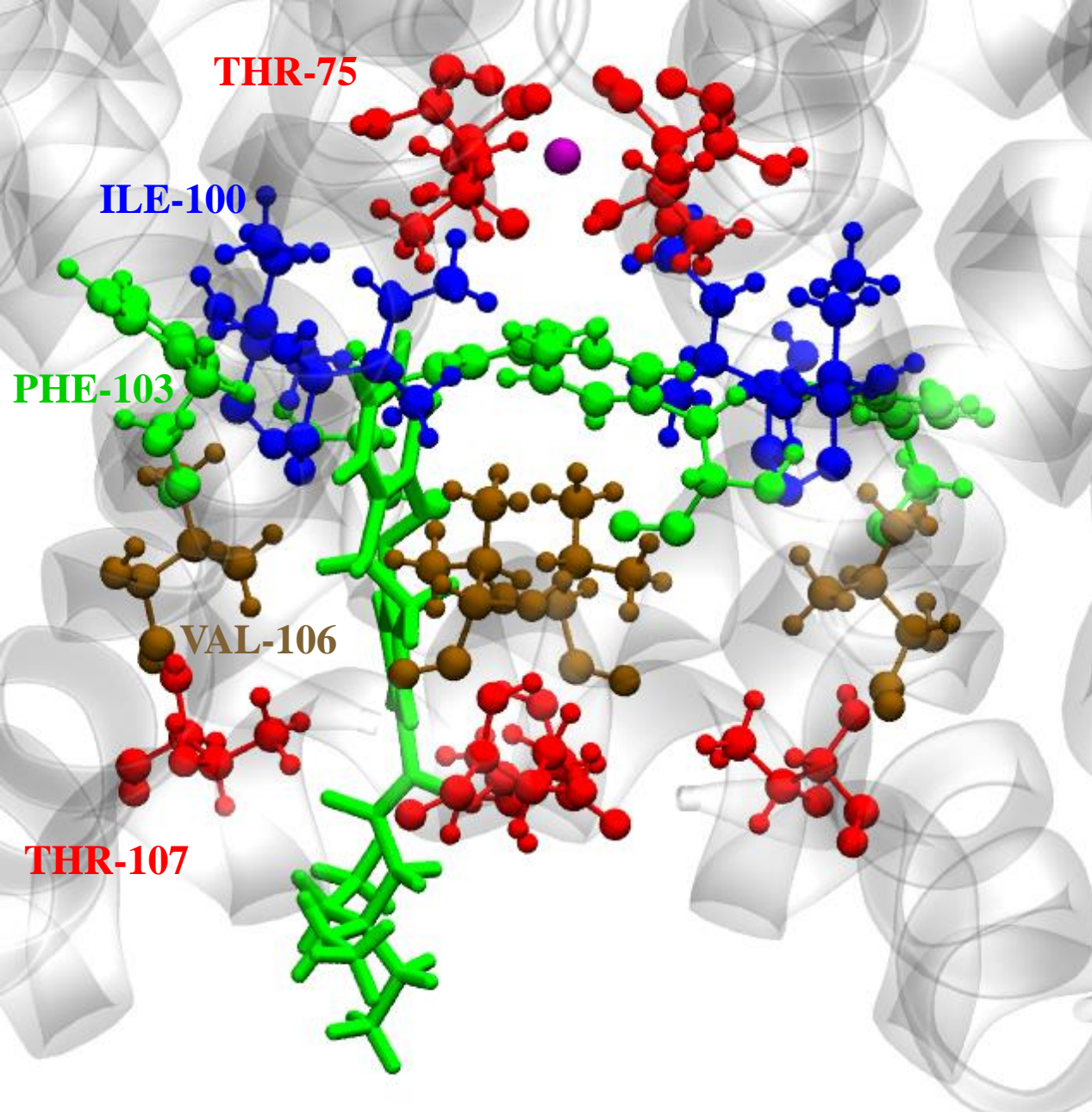




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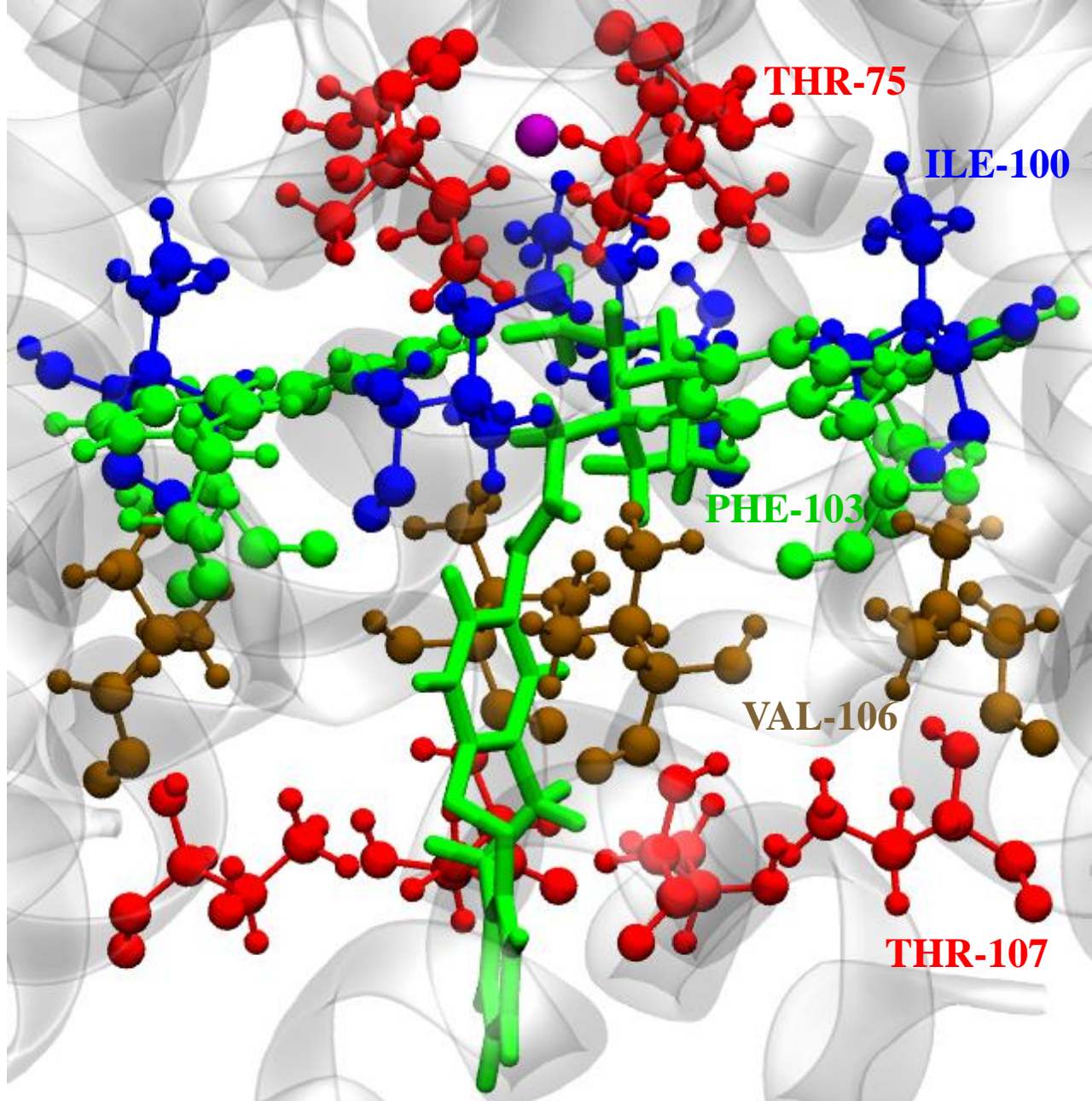


E2 (C0)
MM/PBSA Binding energy = -15.2 kcal/mol³²



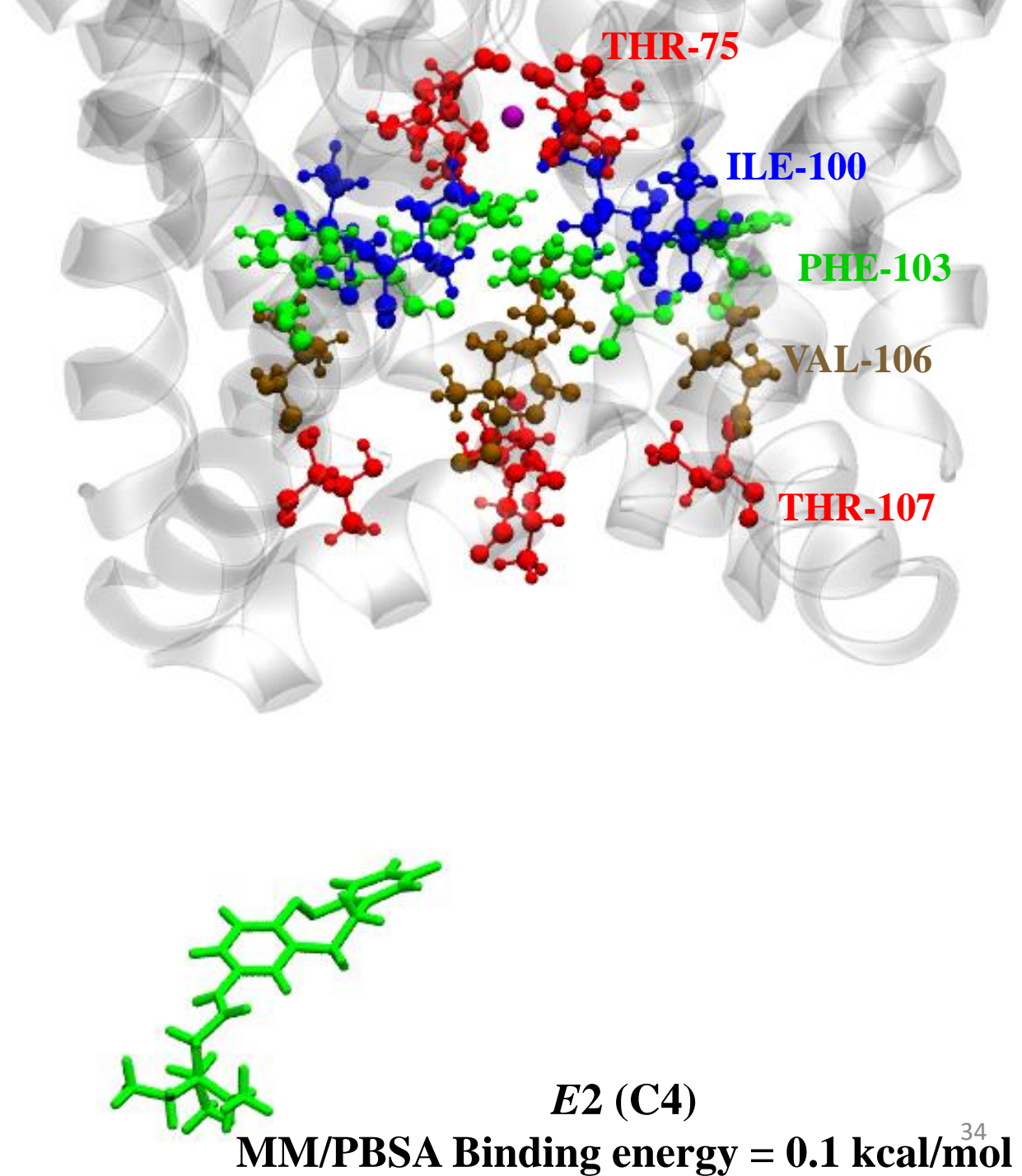
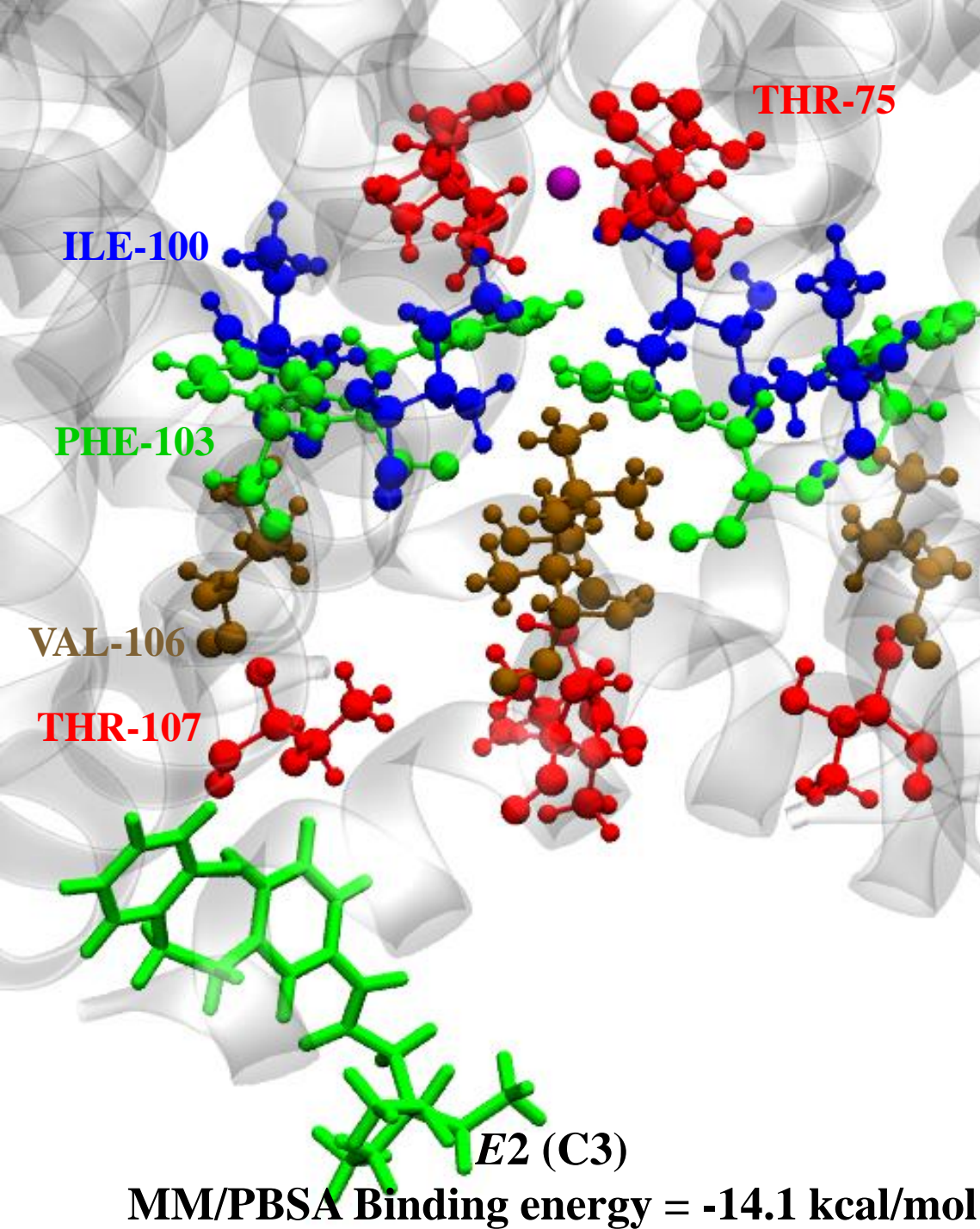
E2 (C1)

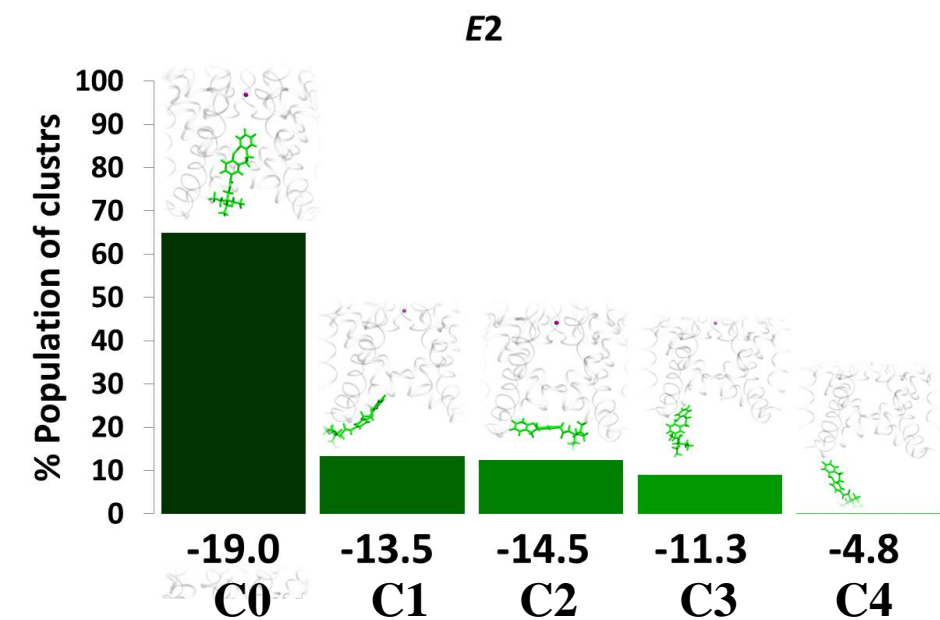
MM/PBSA Binding energy = -19.0 kcal/mol



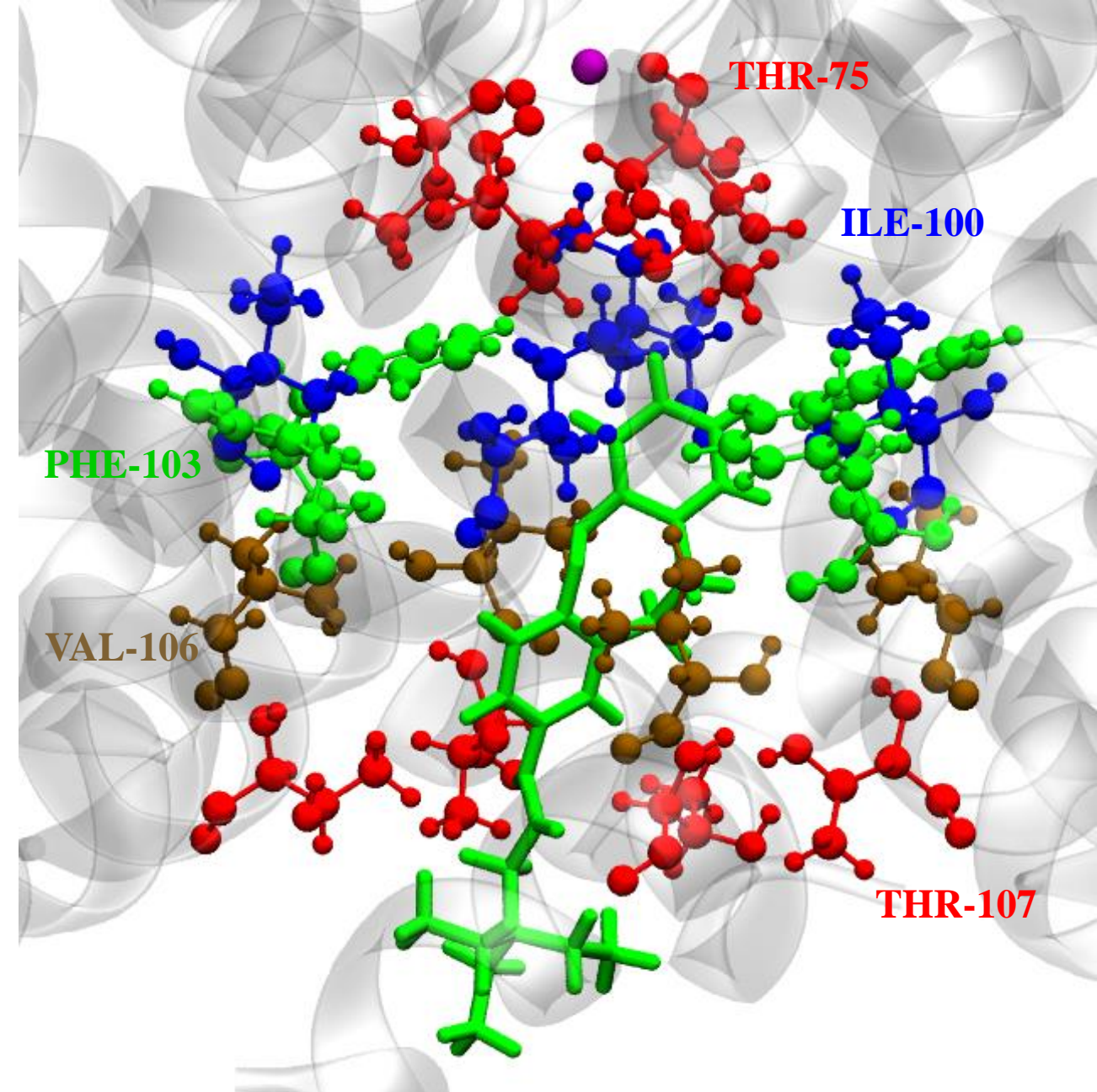
E2 (C2)

MM/PBSA Binding energy = -15.1 kcal/mol³³

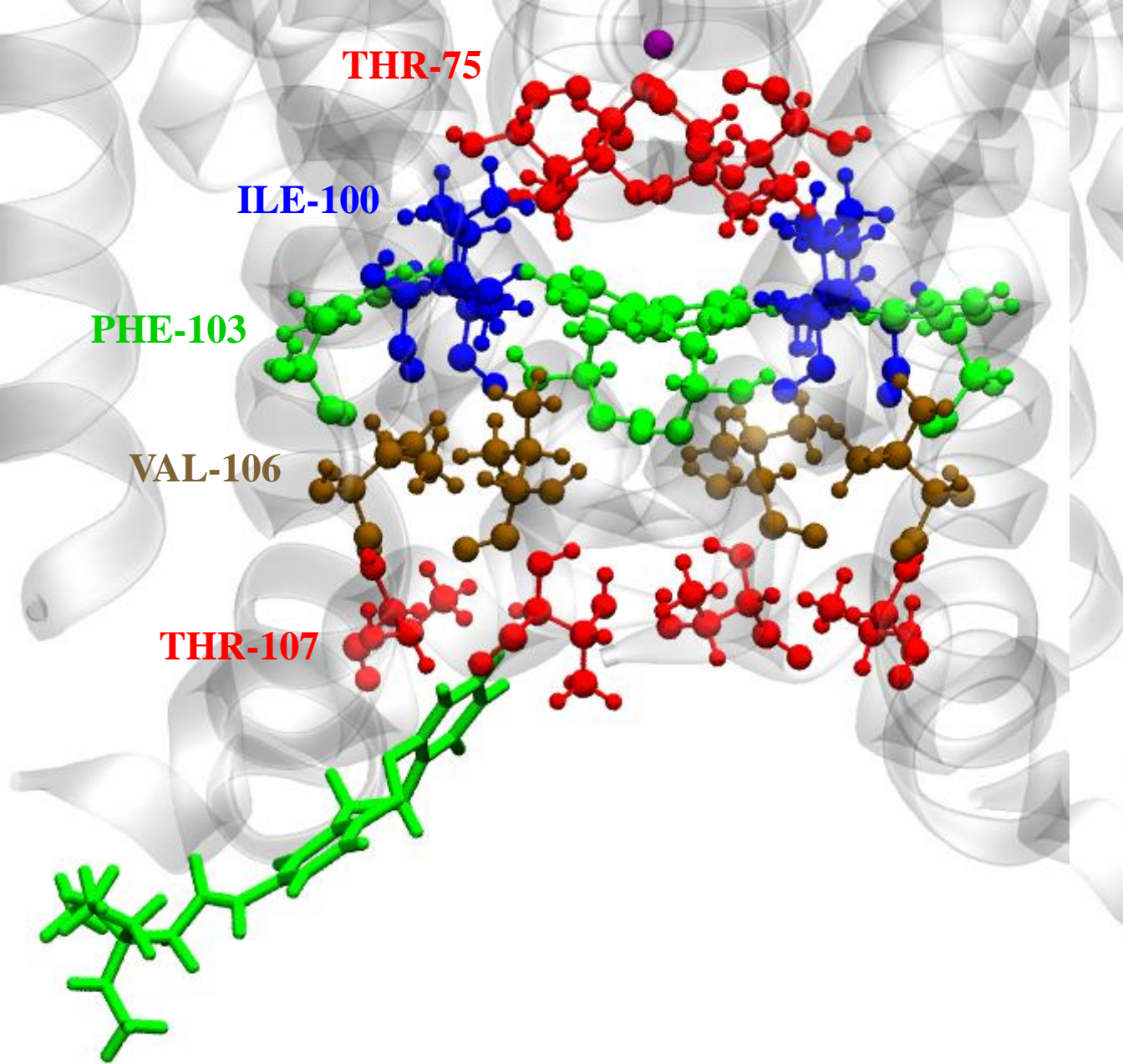




C_n (n = 0 – 1) indicate different clusters. The percentage population of clusters is given on the Y-axis. Corresponding binding energy (B.E.) is given in kcal/mol on the X-axis. Protein is represented in new cartoon model with K⁺-ion (purple ball), and *E*-LAB-TEA is shown in stick model (green colour).

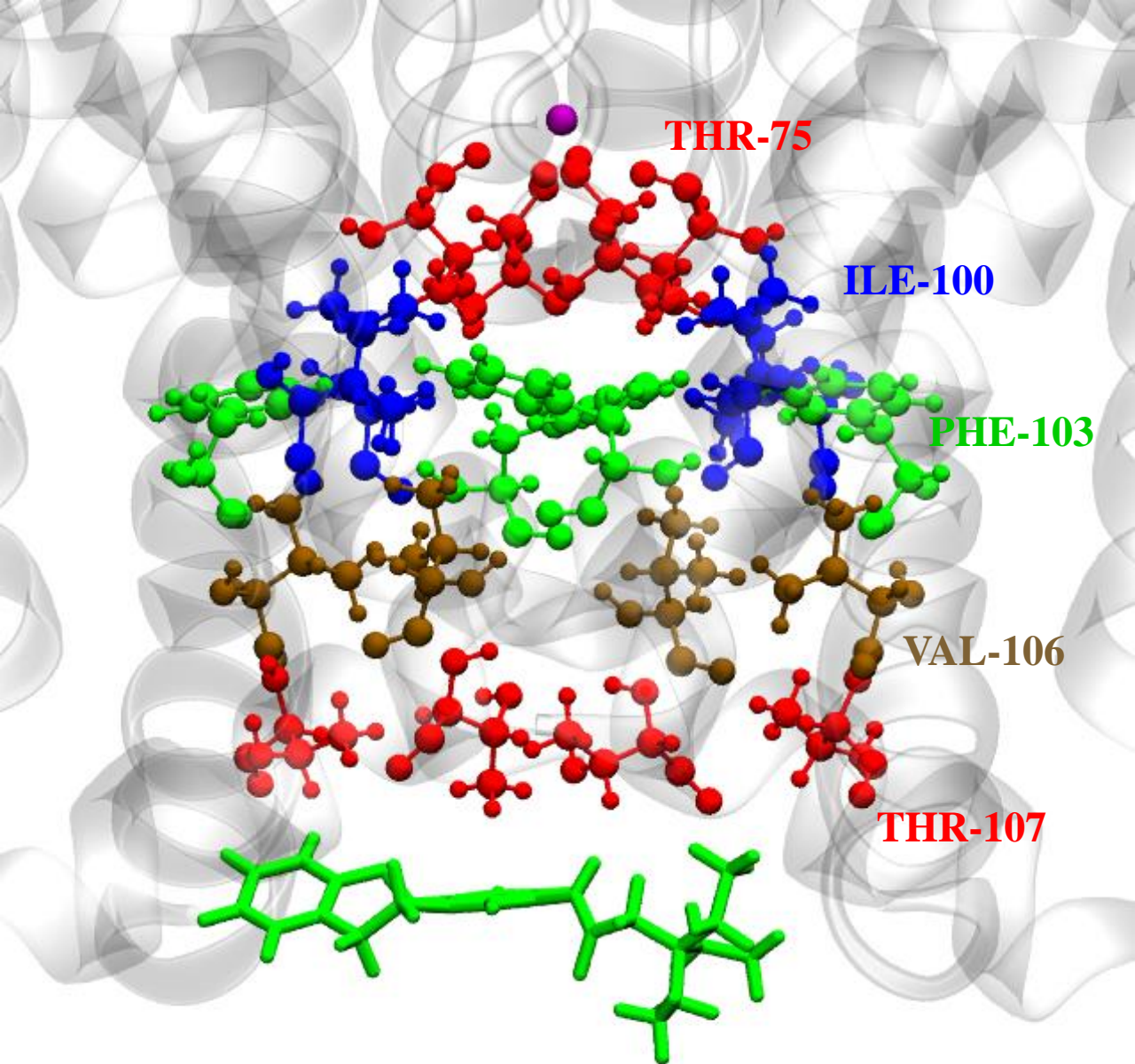


***E*2 (C0)**
MM/PBSA Binding energy = -19.0 kcal/mol³⁵



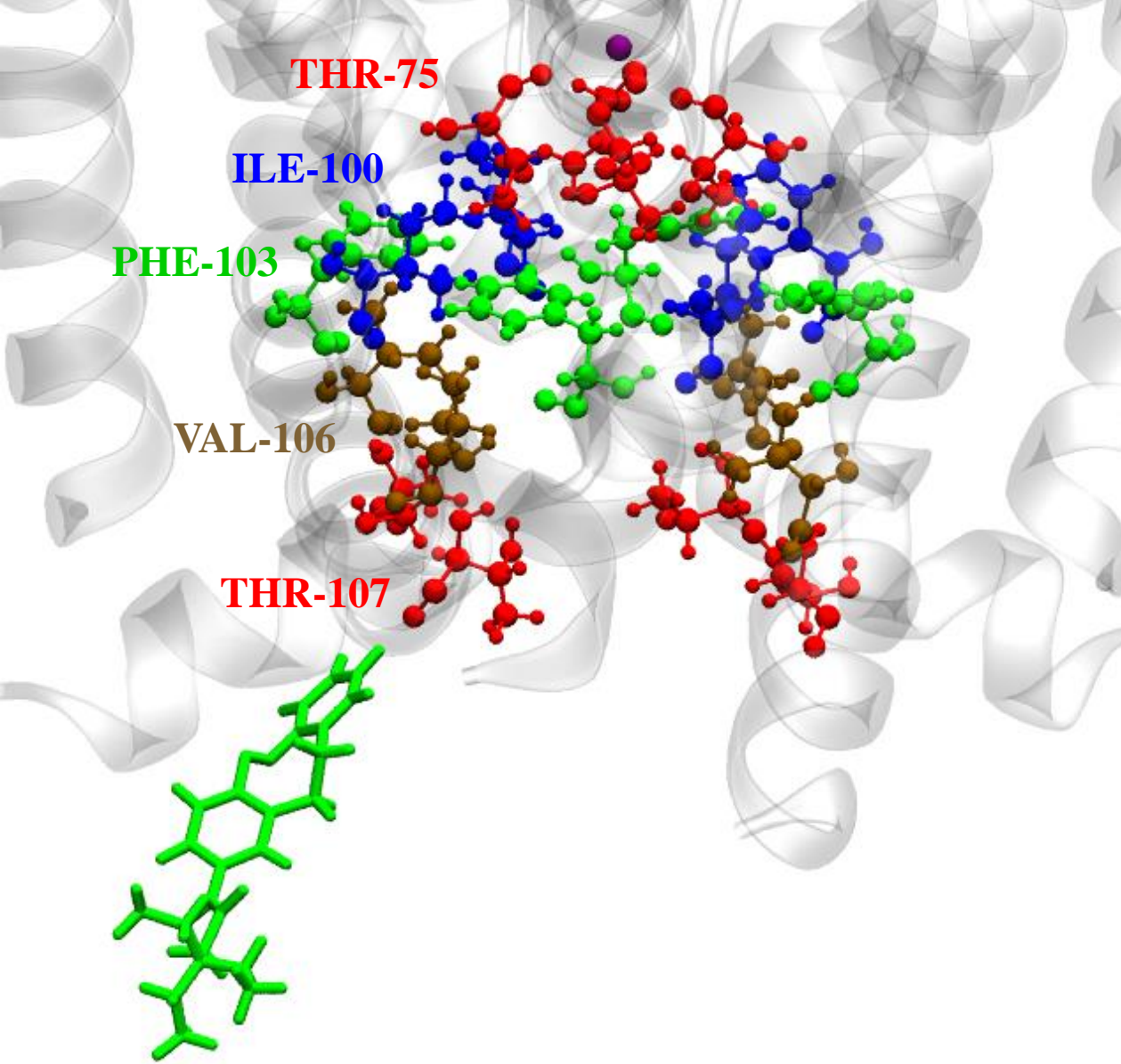
E2 (C1)

MM/PBSA Binding energy = -13.5 kcal/mol



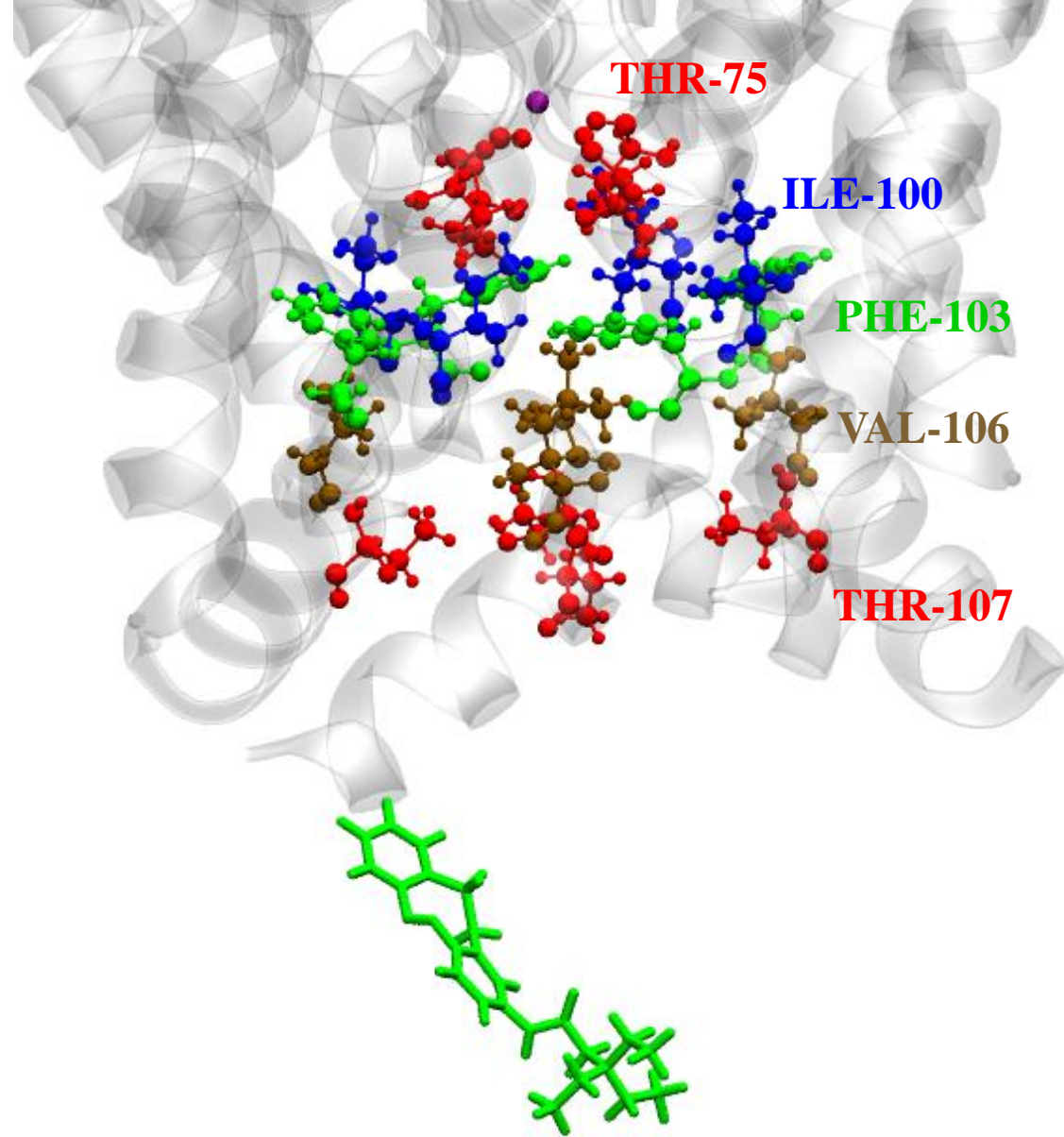
E2 (C2)

MM/PBSA Binding energy = -14.5 kcal/mol³⁶



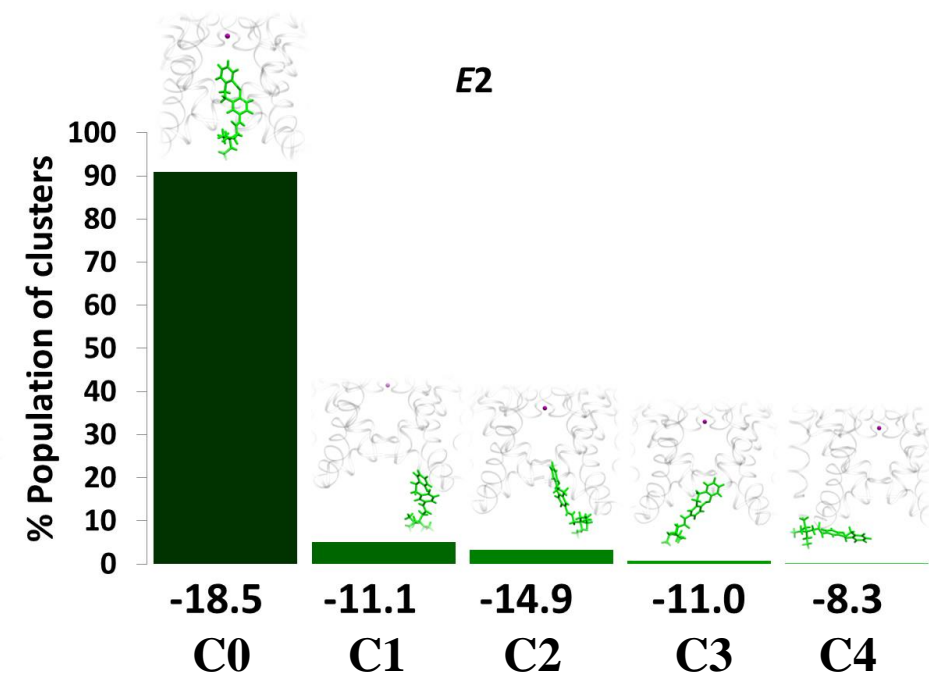
E2 (C3)

MM/PBSA Binding energy = -11.3 kcal/mol

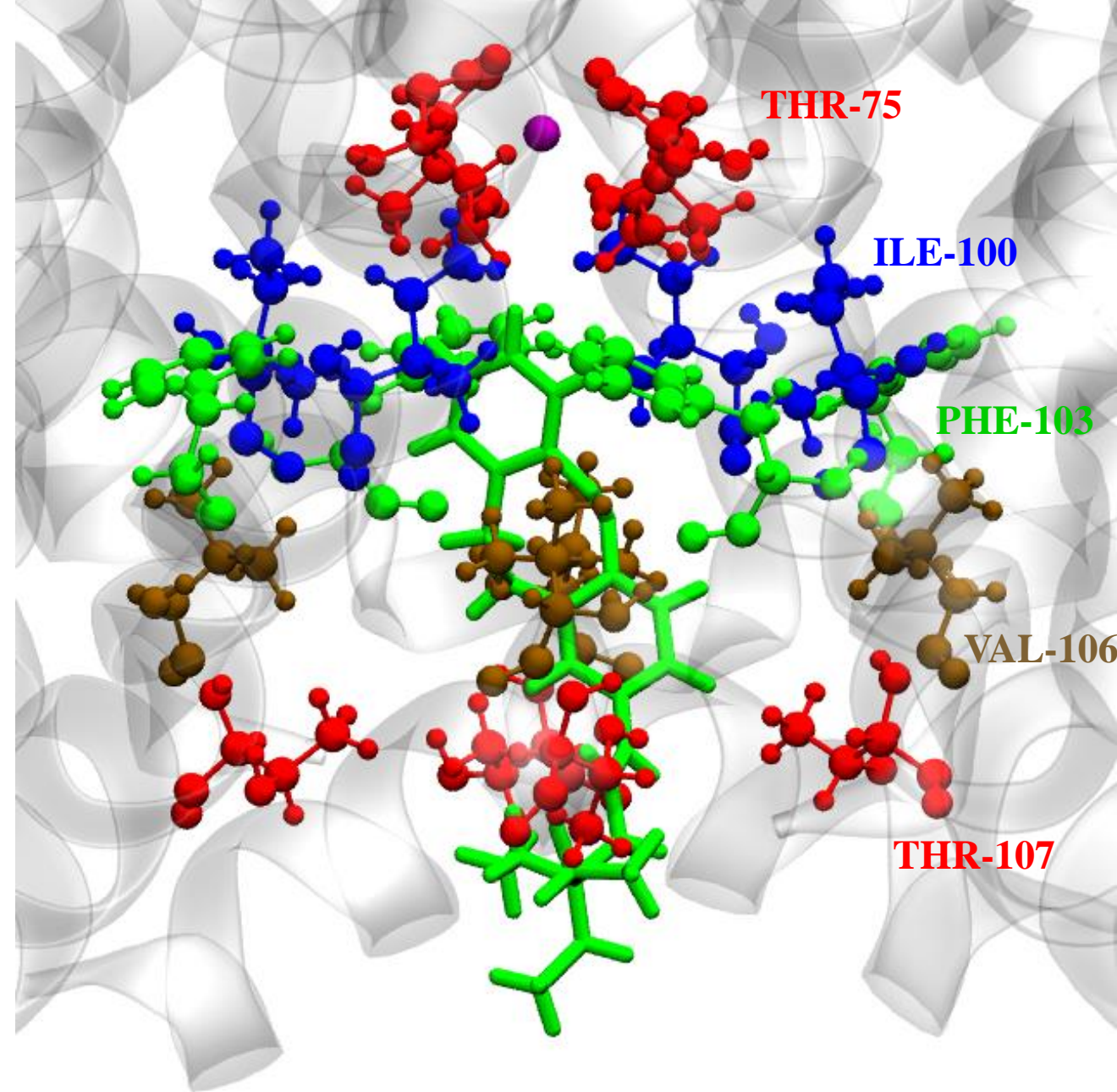


E2 (C4)

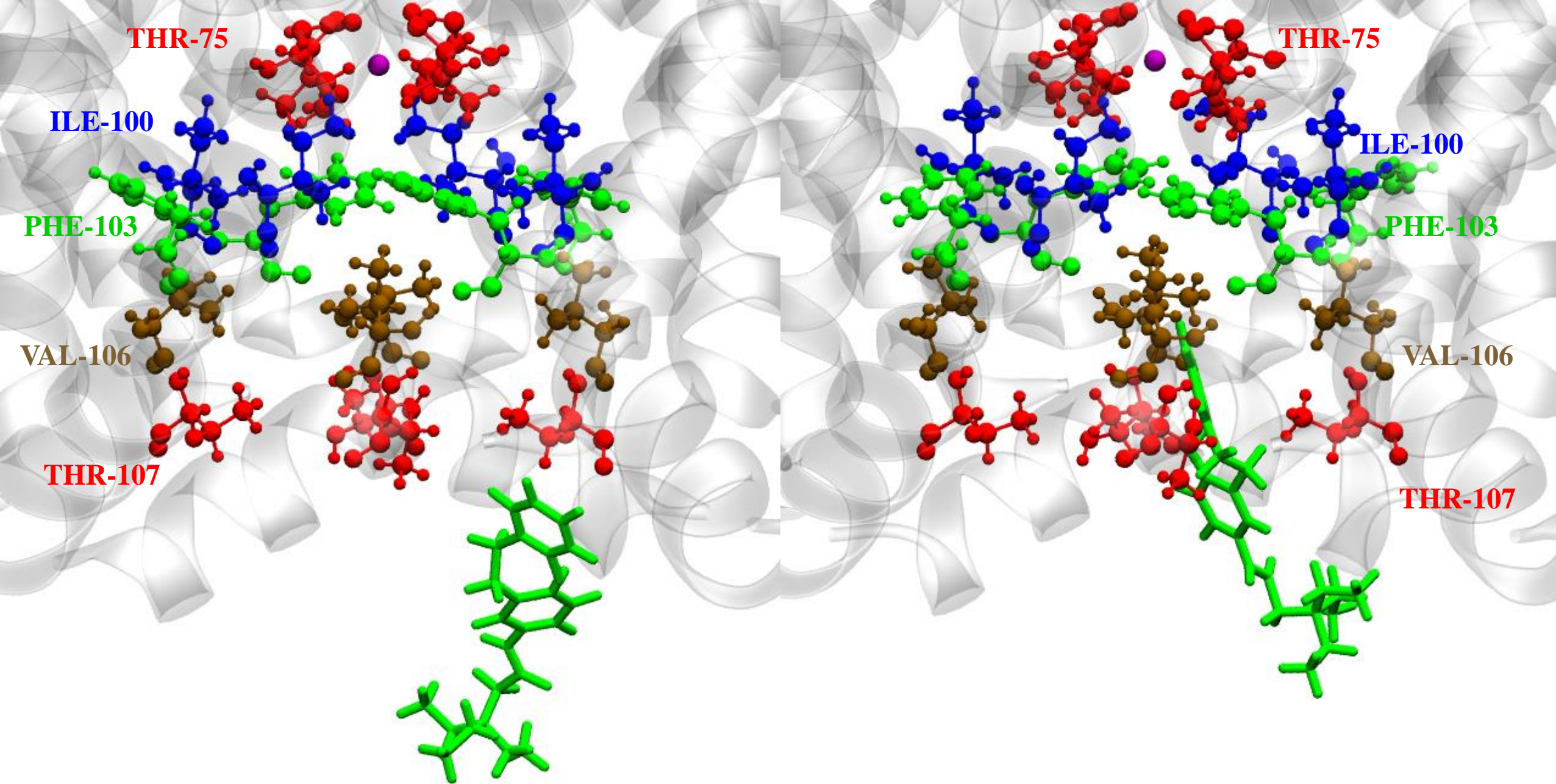
MM/PBSA Binding energy = -4.8 kcal/mol³⁷



C_n (n = 0 – 1) indicate different clusters. The percentage population of clusters is given on the Y-axis. Corresponding binding energy (B.E.) is given in kcal/mol on the X-axis. Protein is represented in new cartoon model with K⁺-ion (purple ball), and *E*-LAB-TEA is shown in stick model (green colour).



***E*2 (C0)**
MM/PBSA Binding energy = -18.5 kcal/mol³⁸

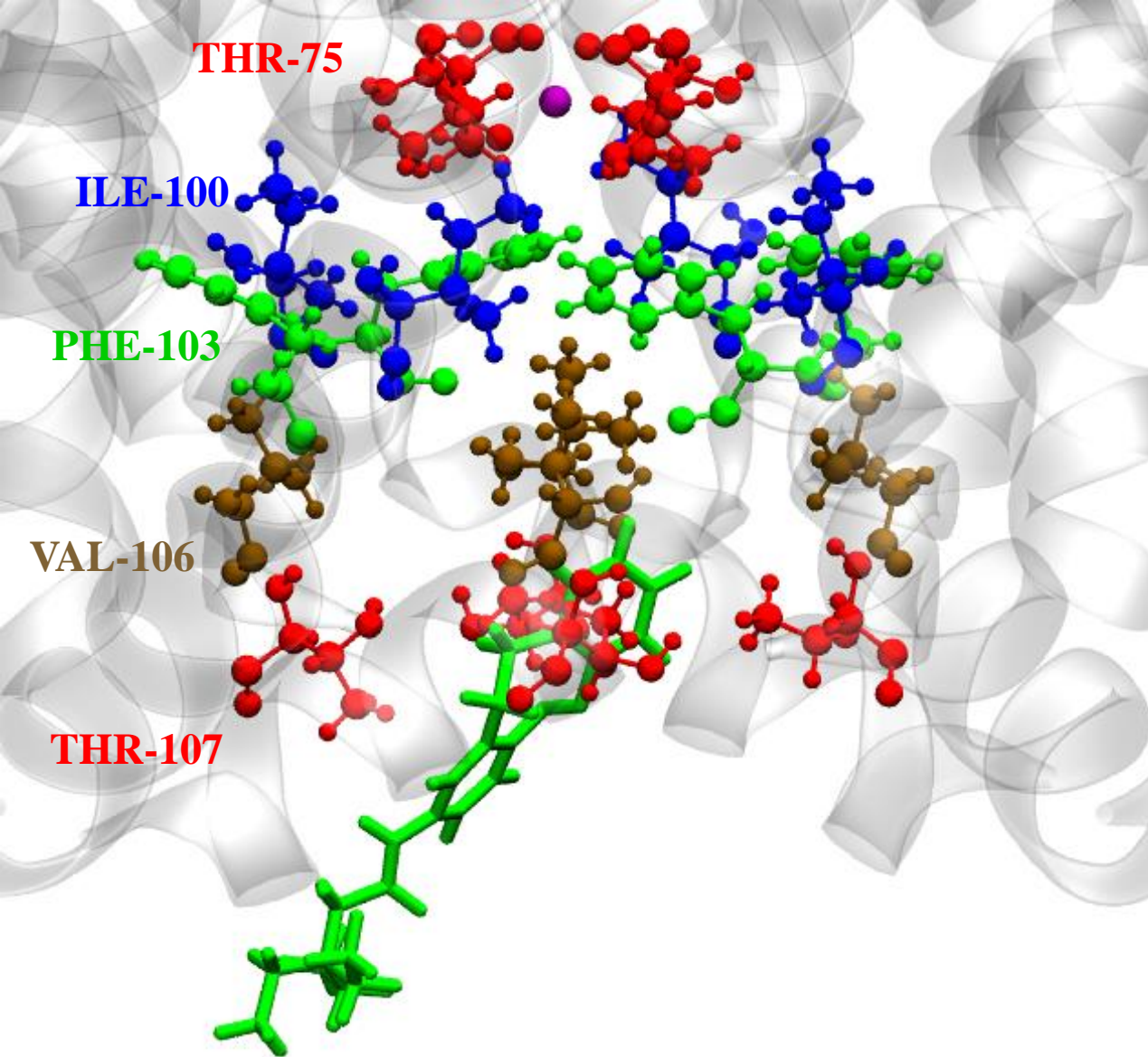


E2 (C1)

MM/PBSA Binding energy = -11.1 kcal/mol

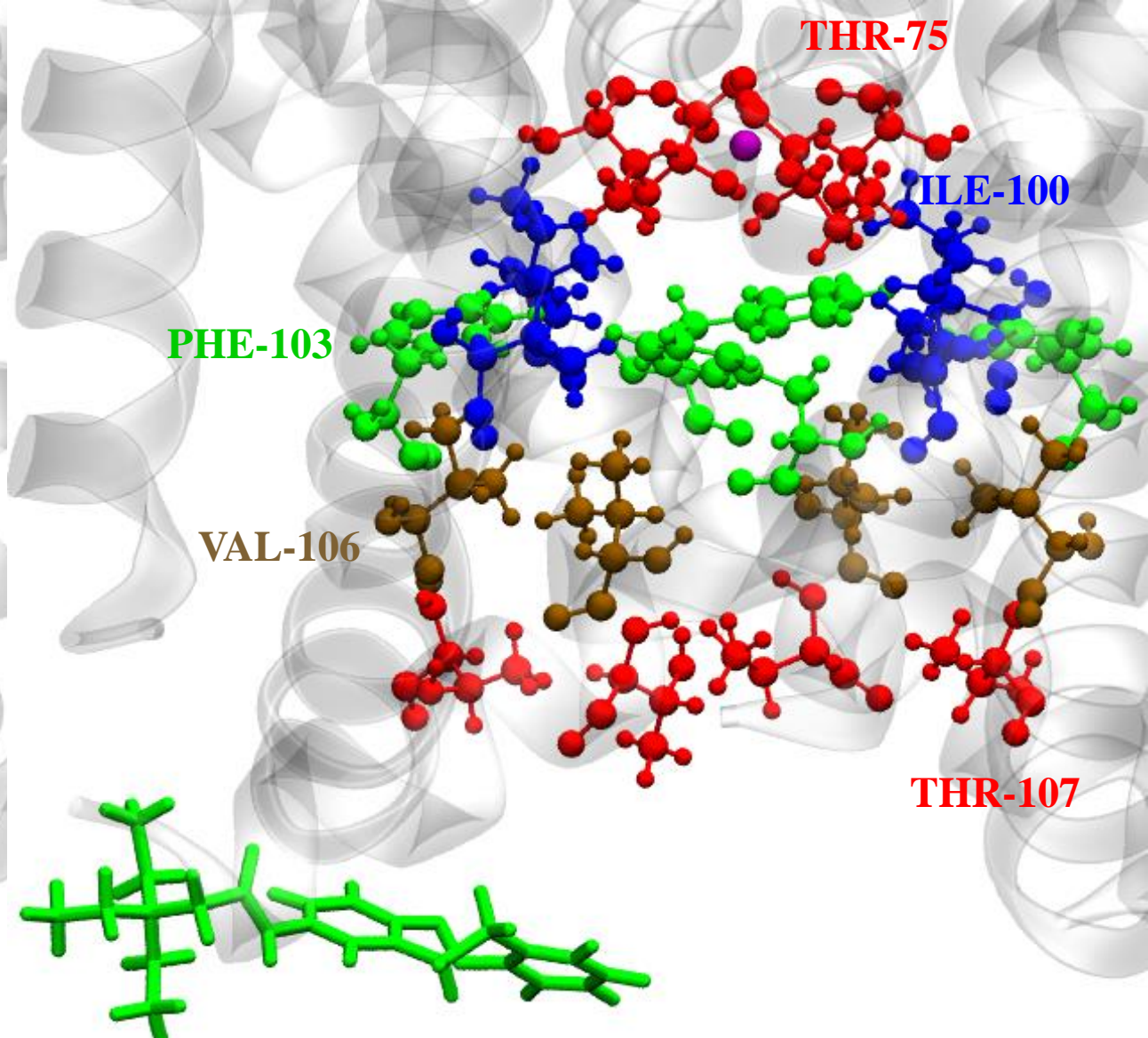
E2 (C2)

MM/PBSA Binding energy = -14.9 kcal/mol³⁹



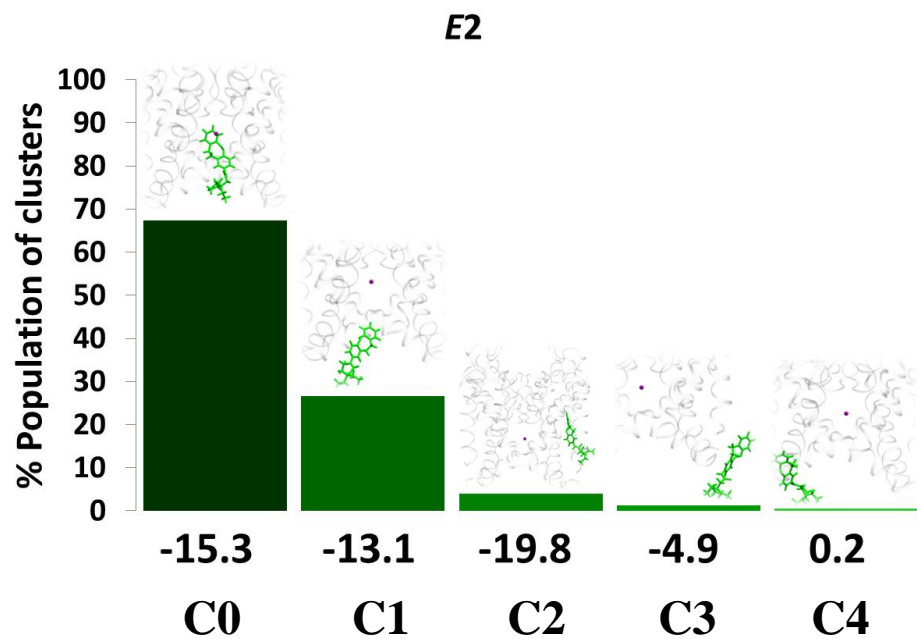
E2 (C3)

MM/PBSA Binding energy = -11.0 kcal/mol

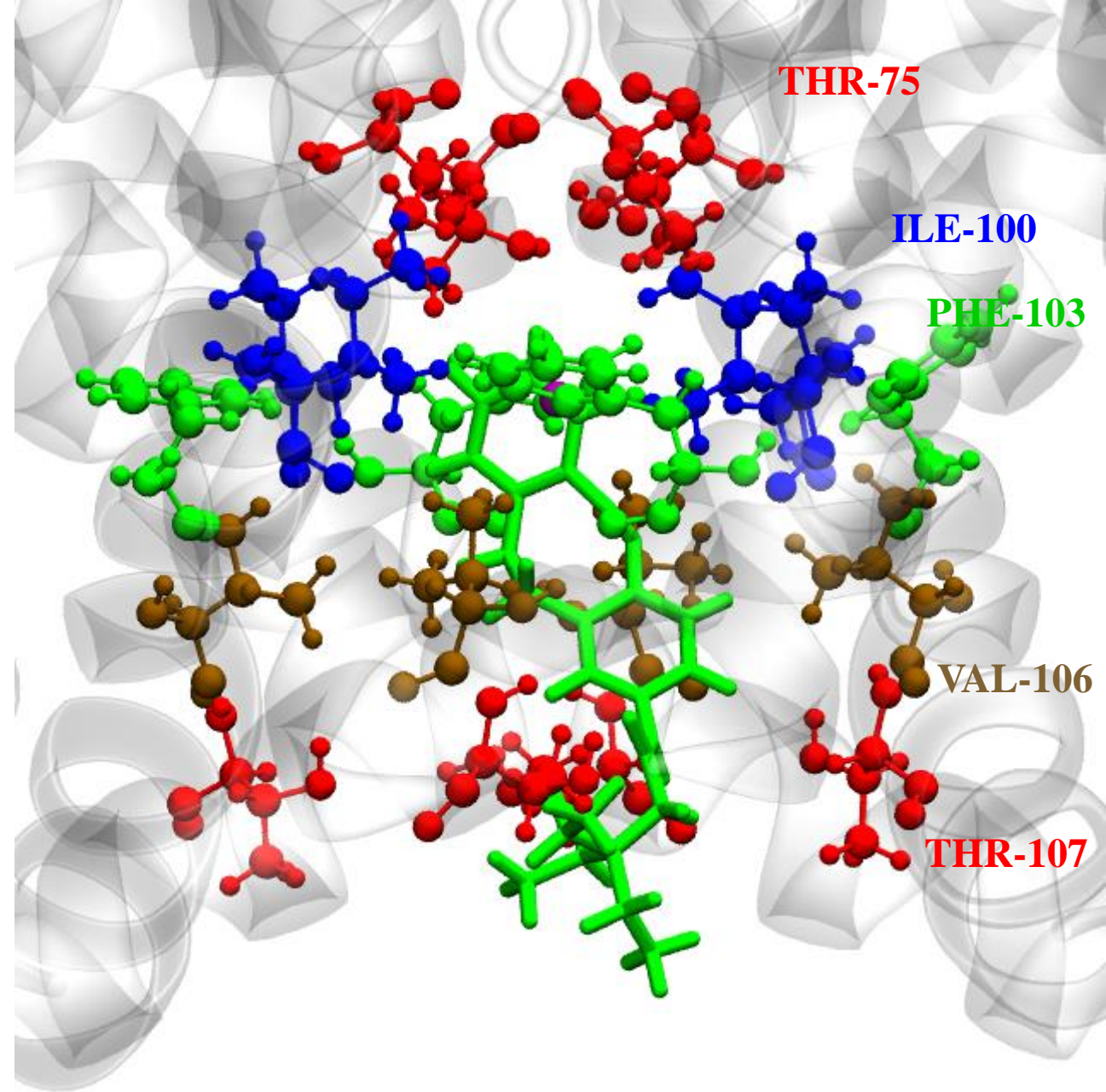


E2 (C4)

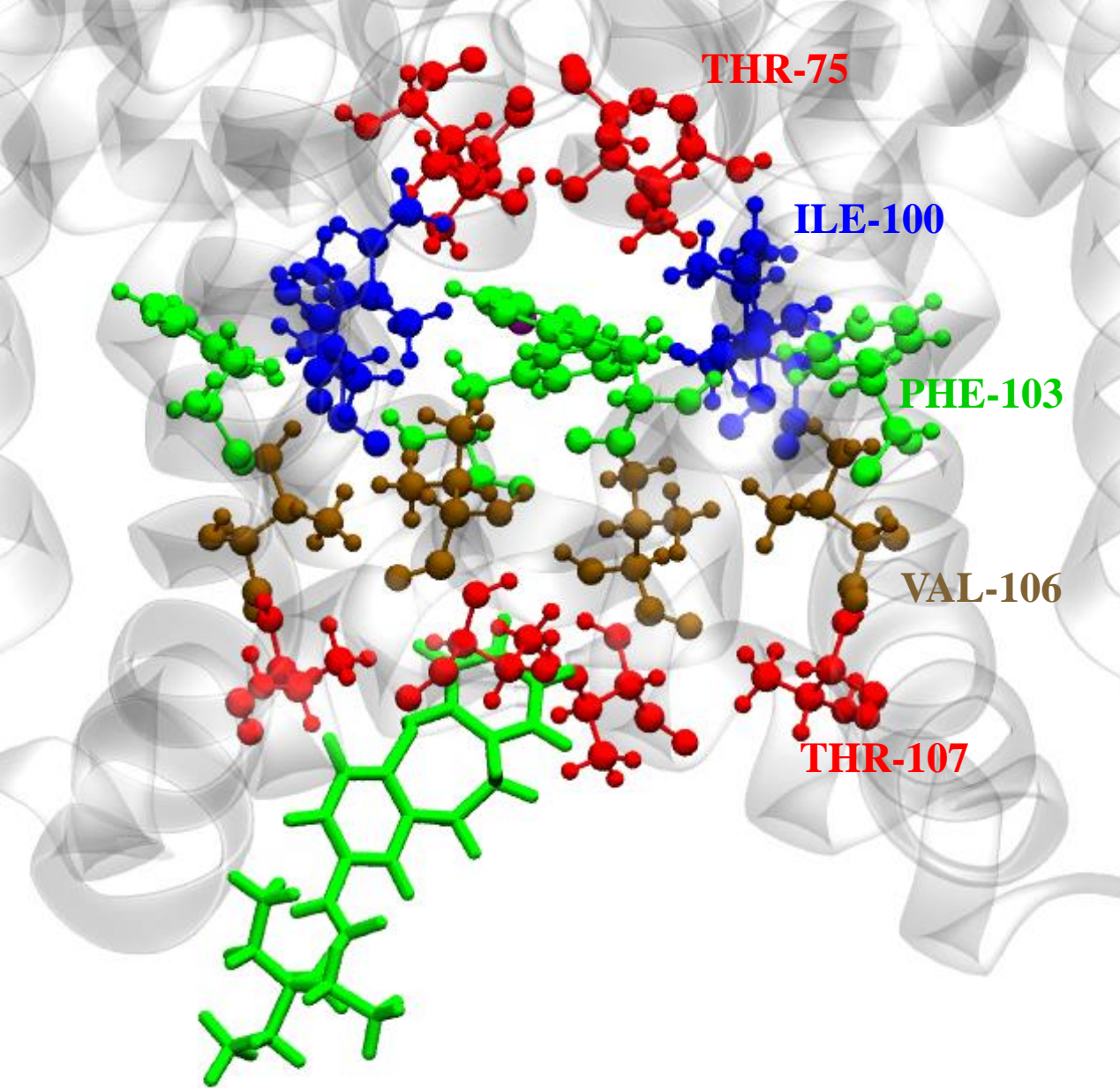
MM/PBSA Binding energy = -8.3 kcal/mol⁴⁰



C_n (n = 0 – 1) indicate different clusters. The percentage population of clusters is given on the Y-axis. Corresponding binding energy (B.E.) is given in kcal/mol on the X-axis. Protein is represented in new cartoon model with K⁺-ion (purple ball), and *E*-LAB-TEA is shown in stick model (green colour).

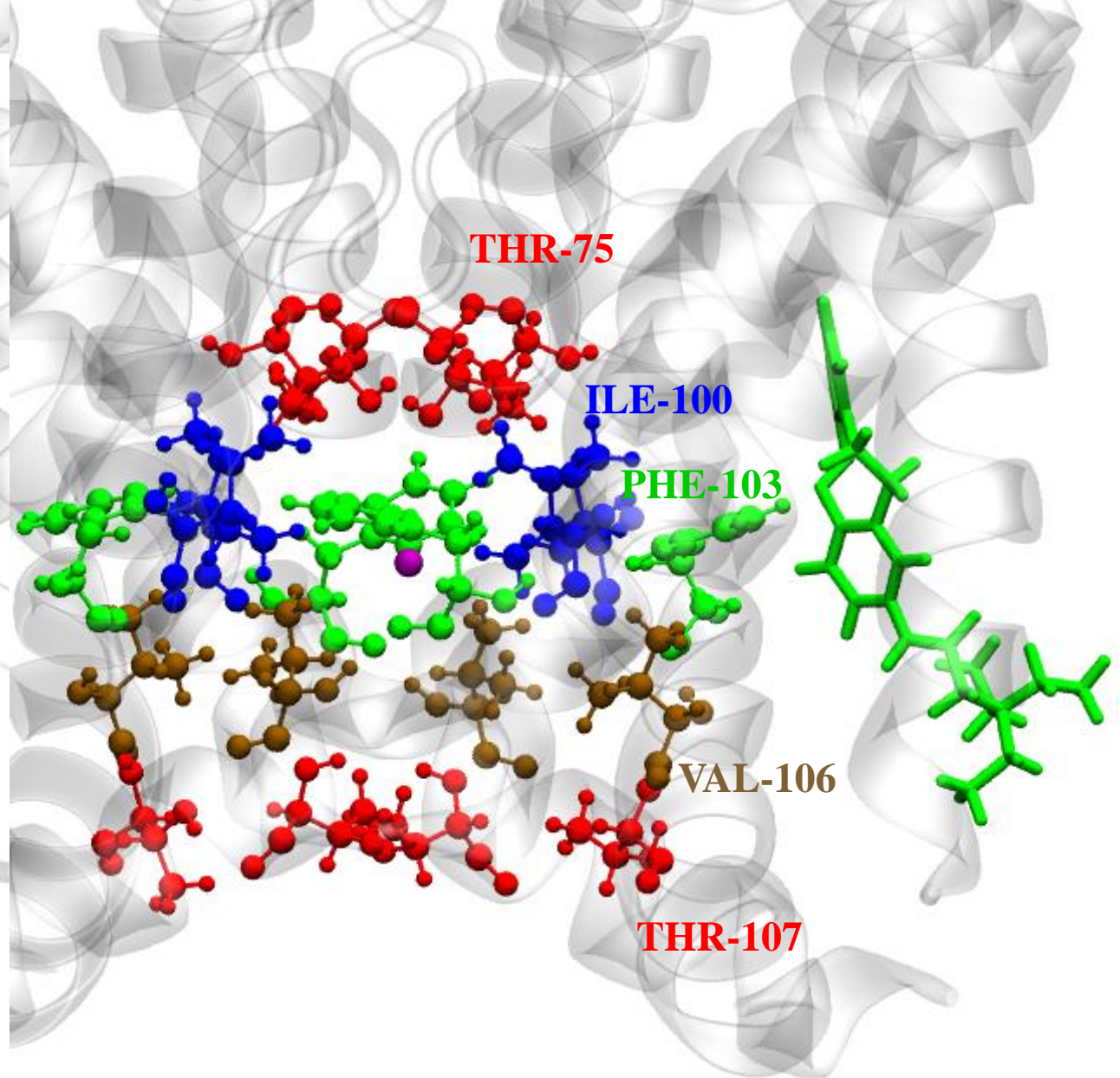


E2 (C0)
MM/PBSA Binding energy = -15.3 kcal/mol⁴¹



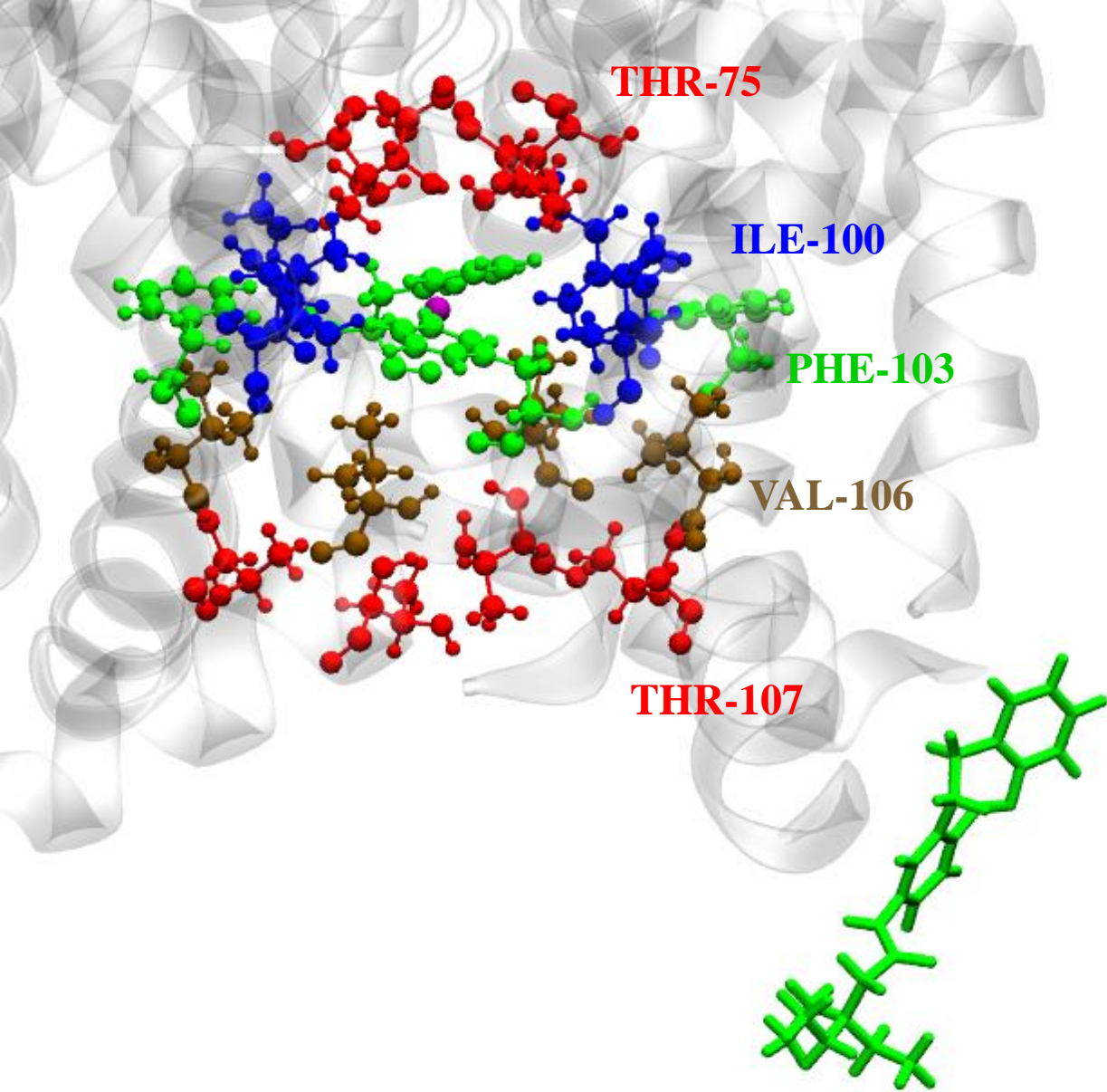
E2 (C1)

MM/PBSA Binding energy = -13.1 kcal/mol



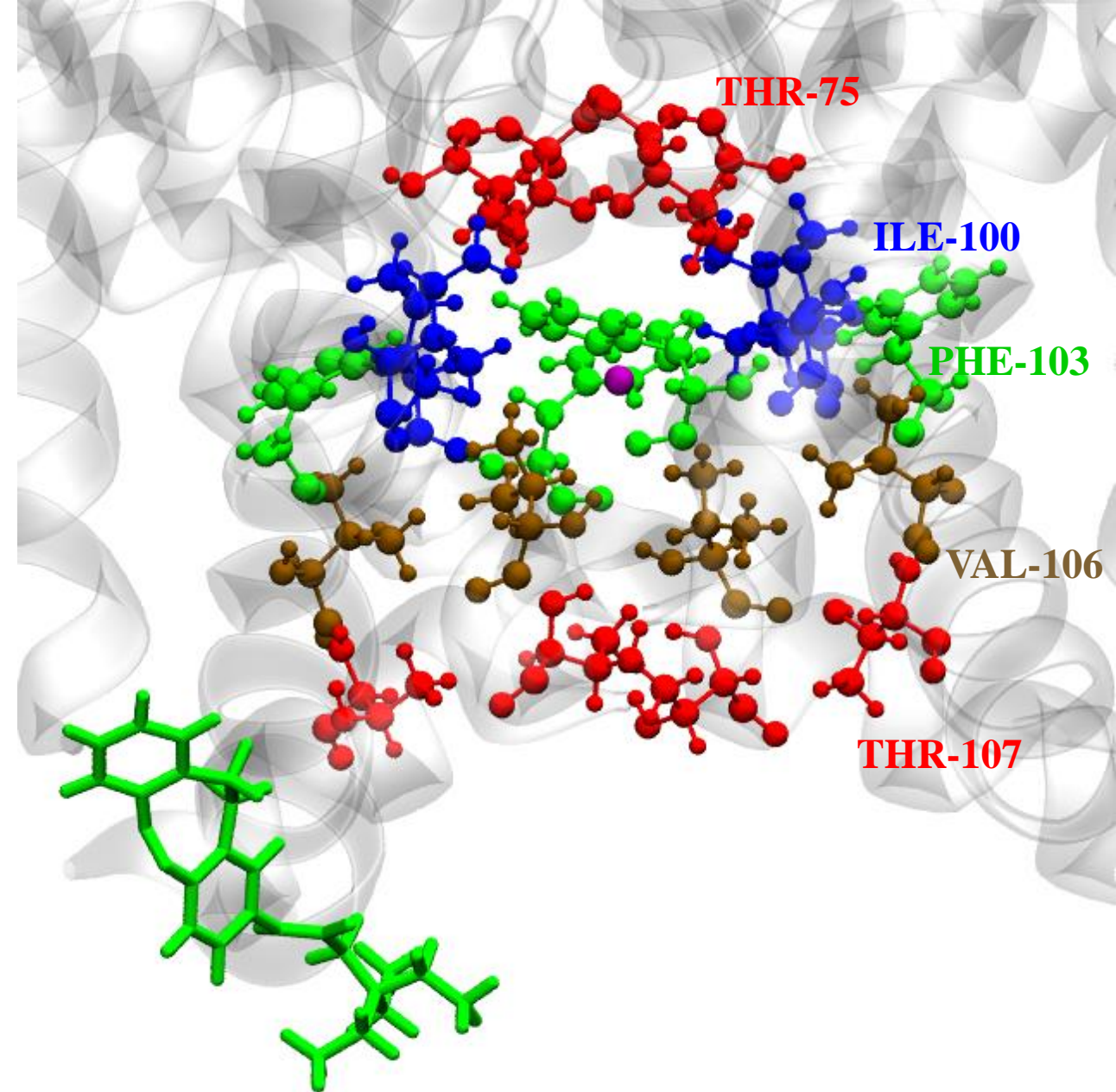
E2 (C2)

MM/PBSA Binding energy = -19.8 kcal/mol⁴²



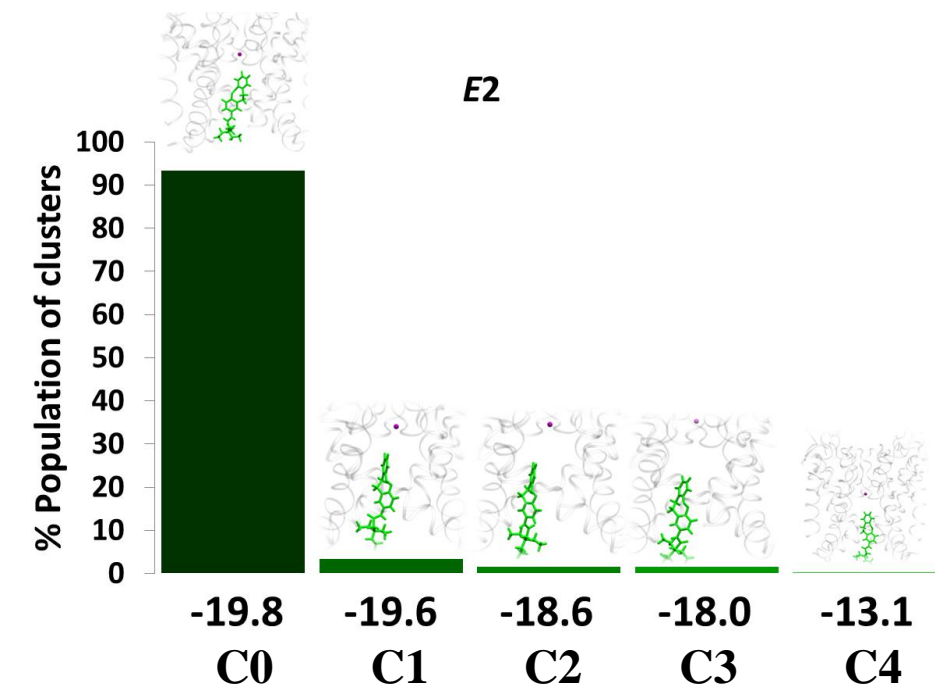
E2 (C3)

MM/PBSA Binding energy = -4.9 kcal/mol

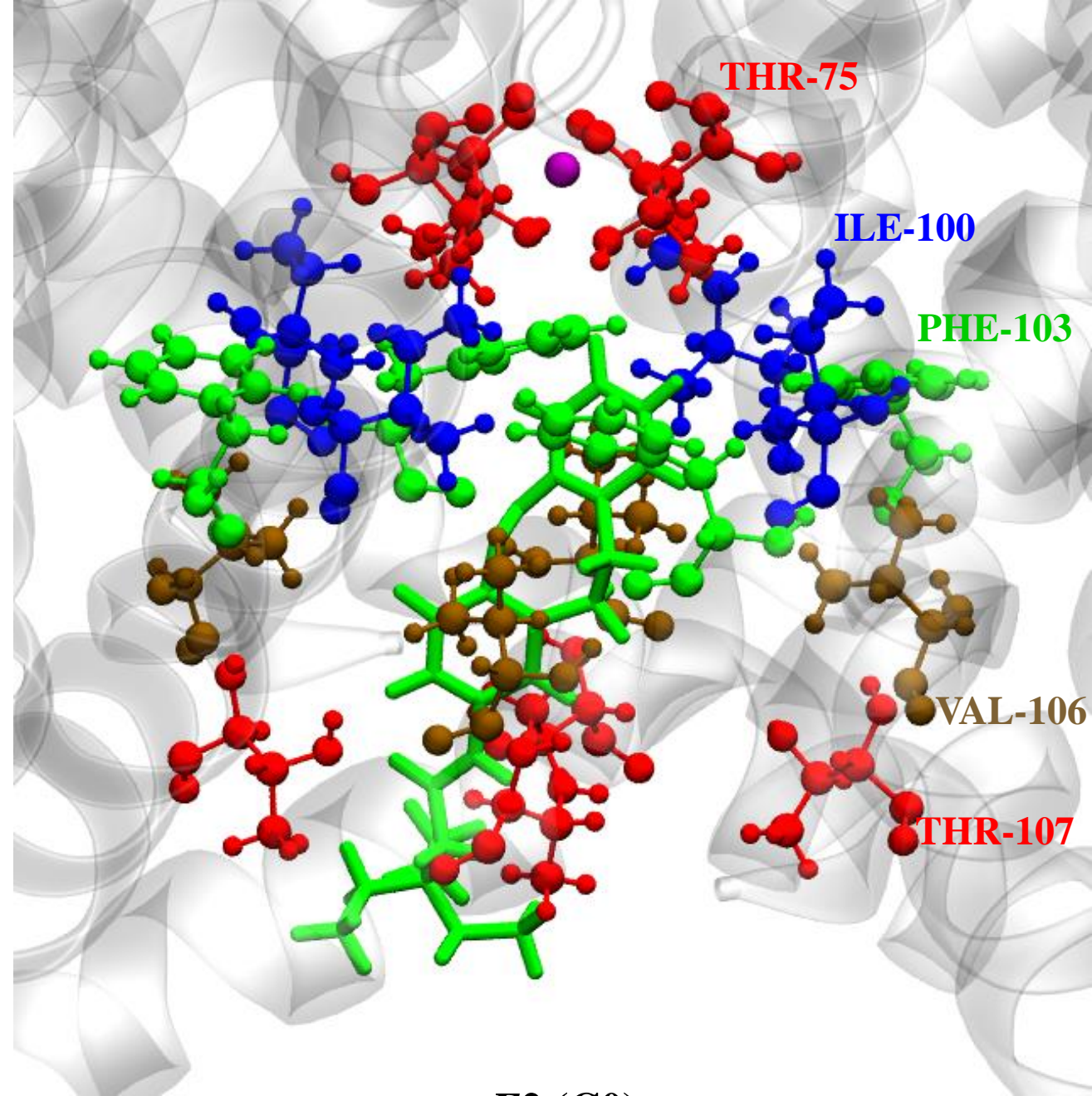


E2 (C4)

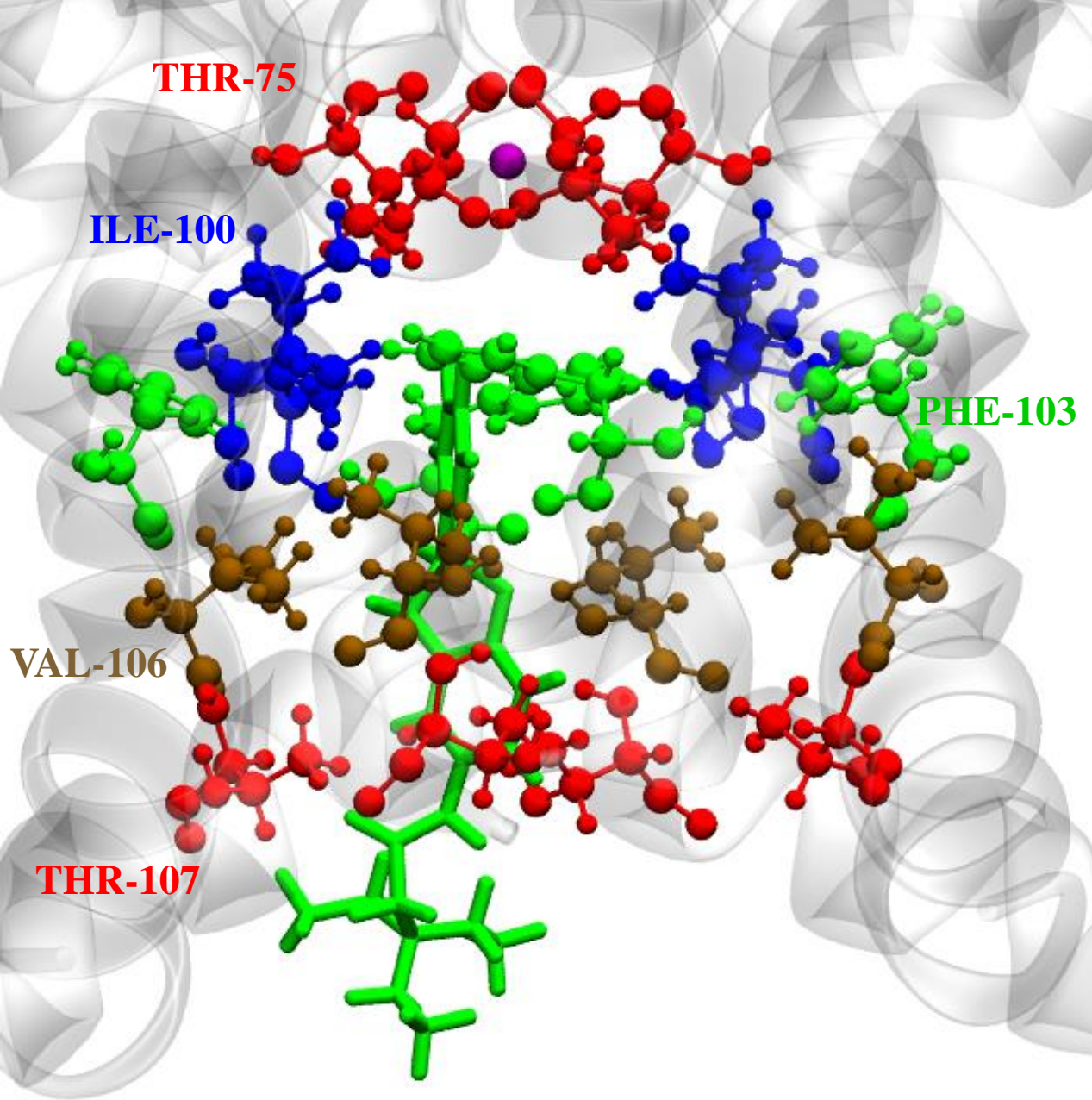
MM/PBSA Binding energy = 0.2 kcal/mol



C_n (n = 0 – 1) indicate different clusters. The percentage population of clusters is given on the Y-axis. Corresponding binding energy (B.E.) is given in kcal/mol on the X-axis. Protein is represented in new cartoon model with K⁺-ion (purple ball), and *E*-LAB-TEA is shown in stick model (green colour).

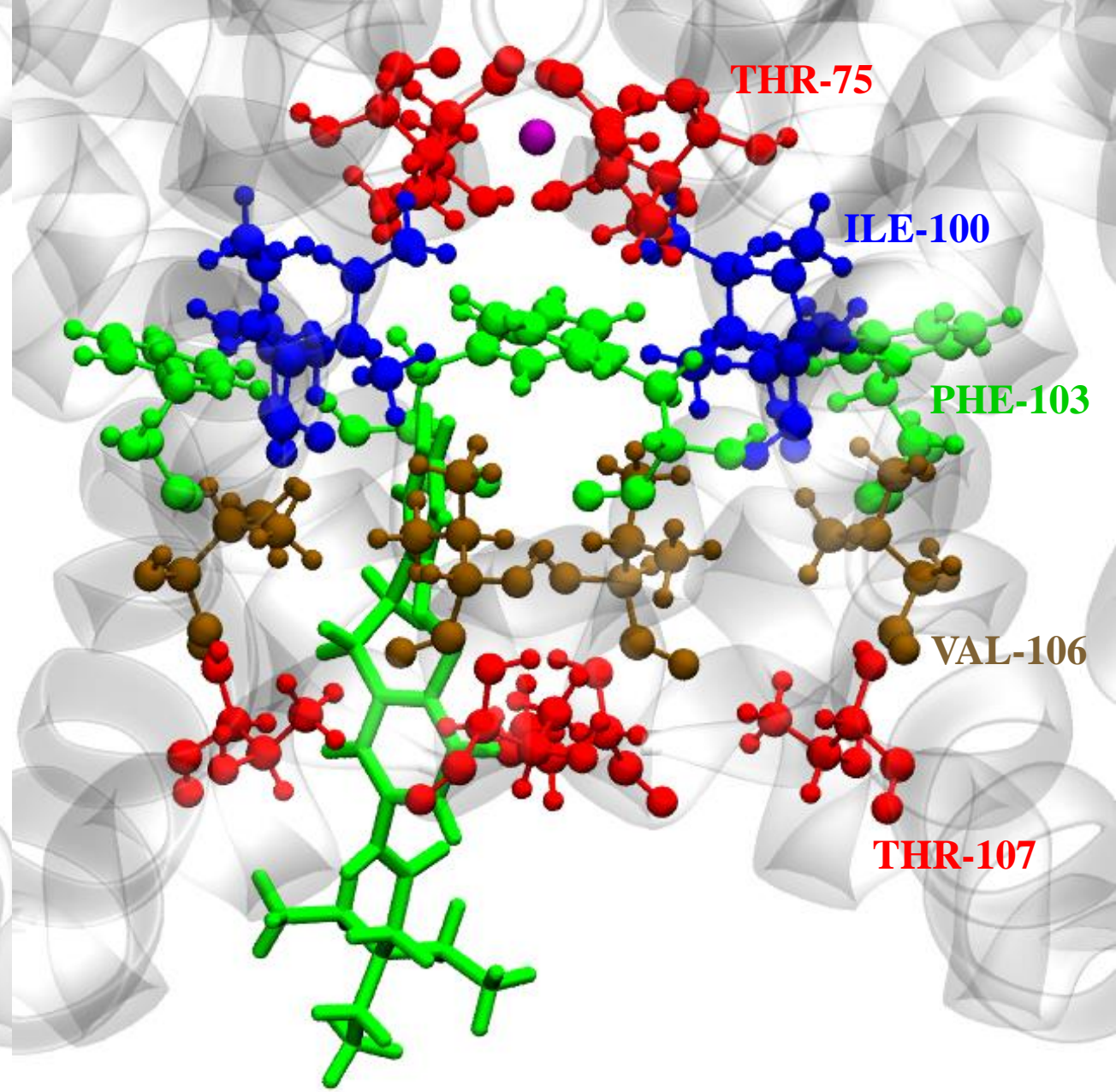


***E*2 (C0)**
MM/PBSA Binding energy = -19.8 kcal/mol



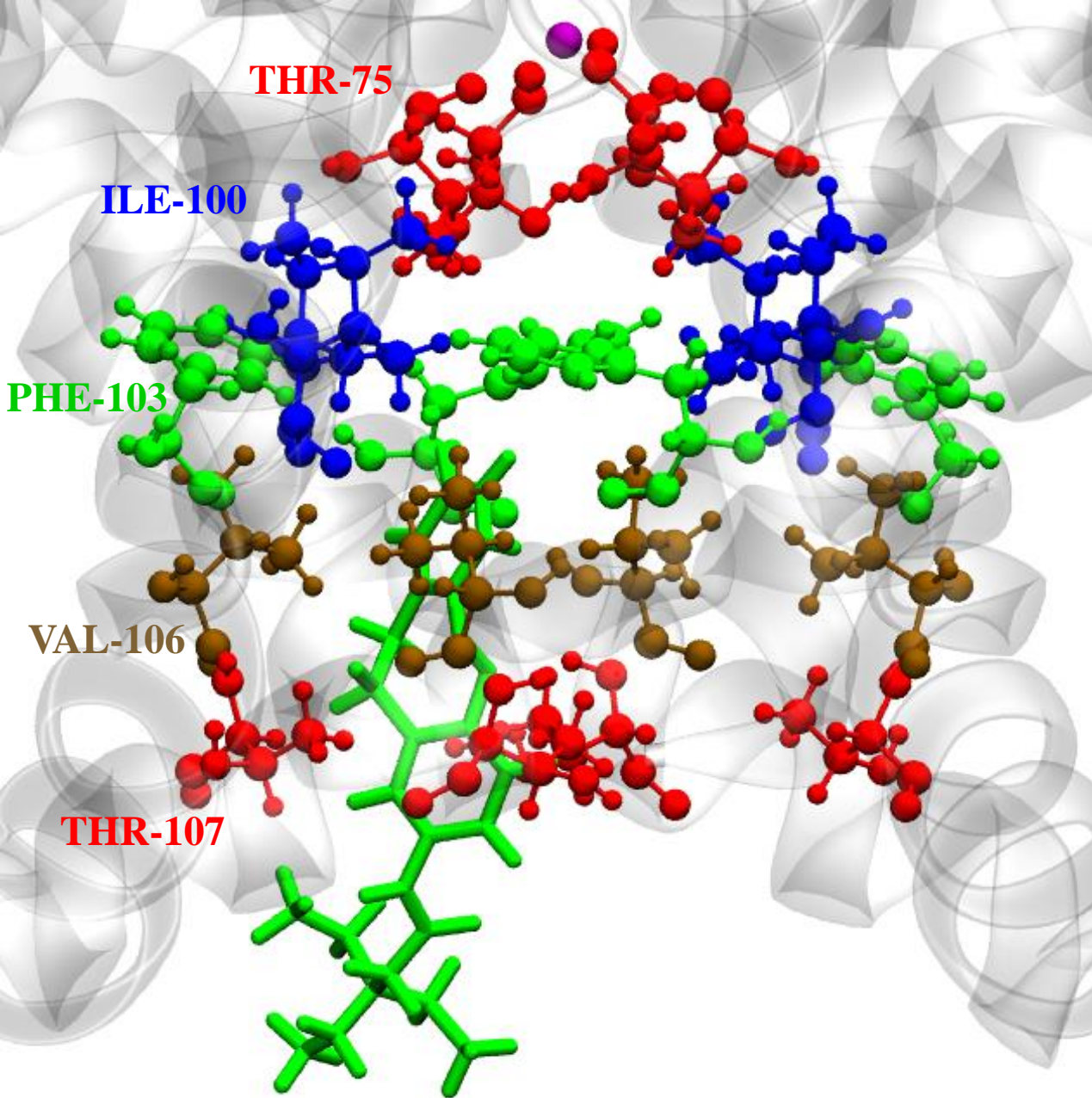
E2 (C1)

MM/PBSA Binding energy = -19.6 kcal/mol



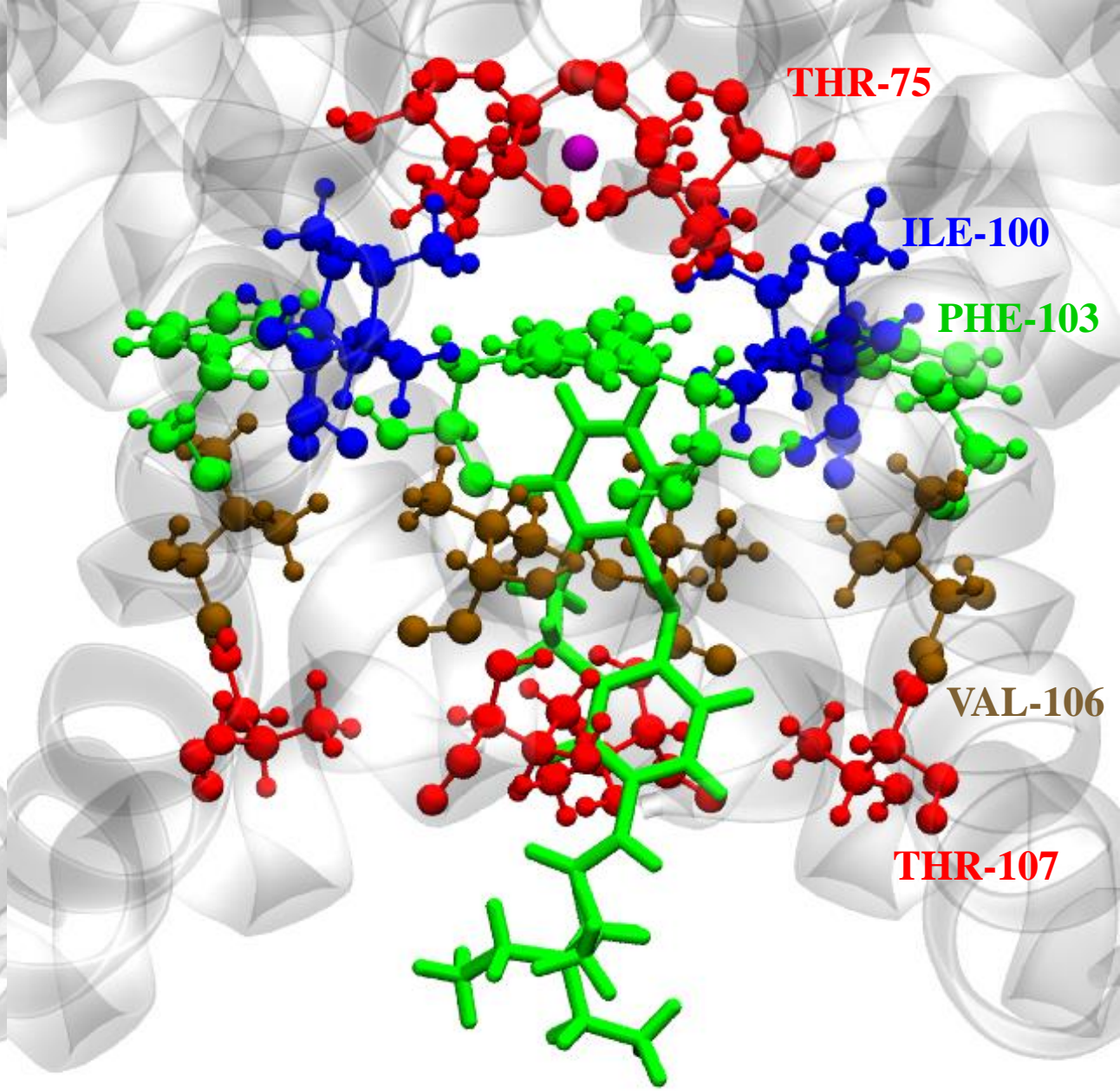
E2 (C2)

MM/PBSA Binding energy = -18.6 kcal/mol⁴⁵



E2 (C3)

MM/PBSA Binding energy = -18.0 kcal/mol



E2 (C4)

MM/PBSA Binding energy = -13.1 kcal/mol⁴⁶