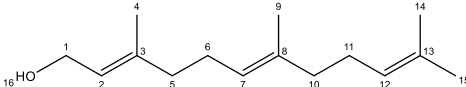


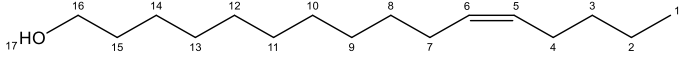
**Table S5.** The distance between the partial carbon atoms on the amino acid residues of OachGOBP1 and each farnesol atom.

OachGOBP1 Farnesol atom	THR32-CA	THR32-CB	THR32-CG2	TRP60-CD1	TRP60-CE2	TRP60-CZ2	Structure
<b>C1</b>	5.1 Å	4.7 Å	4.8 Å	5.0 Å	4.1 Å	3.8 Å	
<b>C2</b>	6.0 Å	5.4 Å	5.7 Å	6.1 Å	5.3 Å	5.0 Å	
<b>C3</b>	6.4 Å	6.4 Å	6.4 Å	7.3 Å	6.6 Å	6.2 Å	
<b>C4</b>	5.8 Å	6.2 Å	6.4 Å	7.9 Å	7.1 Å	6.7 Å	
<b>C5</b>	7.6 Å	7.8 Å	7.6 Å	8.5 Å	7.8 Å	7.5 Å	
<b>C6</b>	8.8 Å	8.7 Å	8.5 Å	8.7 Å	7.9 Å	7.5 Å	
<b>C7</b>	10.0 Å	10.0 Å	9.8 Å	10.1 Å	9.2 Å	8.6 Å	
<b>C8</b>	10.4 Å	10.5 Å	10.5 Å	10.6 Å	9.4 Å	8.6 Å	
<b>C9</b>	9.8 Å	9.8 Å	10.0 Å	9.9 Å	8.4 Å	7.5 Å	
<b>C10</b>	11.7 Å	11.9 Å	11.9 Å	12.1 Å	10.9 Å	10.0 Å	
<b>C11</b>	11.8 Å	12.2 Å	12.3 Å	12.9 Å	11.8 Å	11.0 Å	
<b>C12</b>	11.5 Å	12.0 Å	12.3 Å	13.1 Å	11.8 Å	10.9 Å	
<b>C13</b>	10.5 Å	11.0 Å	11.4 Å	12.5 Å	11.3 Å	10.5 Å	
<b>C14</b>	9.3 Å	9.9 Å	10.2 Å	11.5 Å	10.5 Å	9.8 Å	
<b>C15</b>	10.5 Å	11.2 Å	11.7 Å	13.0 Å	11.7 Å	10.8 Å	
<b>O16</b>	4.4 Å	3.7 Å	3.5 Å	4.0 Å	3.7 Å	3.8 Å	

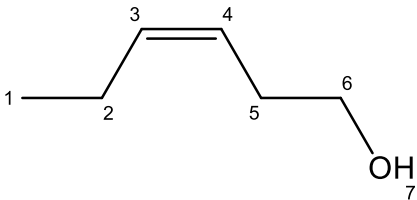
Note: “partial carbon atoms” refers to the carbon atom on an amino acid hydrogen-bonded to the ligand and within the range of ligand 4 Å.

The atoms on all ligands (except hydrogen atoms) are numbered for statistical purposes. The atom numbers are shown in the structure column.

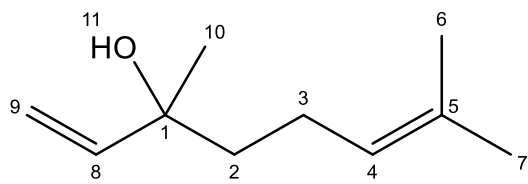
**Table S6.** The distance between the partial carbon atoms on the amino acid residues of OachGOBP1 and each Z11-16: OH atom.

OachGOBP1 Z11-16: OH atom	TRP60-CD1	TRP60-CE2	TRP60-CZ2	TRP60-CH2	Structure
C1	8.3 Å	7.6 Å	7.1 Å	8.1 Å	
C2	9.1 Å	8.2 Å	7.5 Å	8.3 Å	
C3	10.5 Å	9.6 Å	9 Å	9.7 Å	
C4	11.0 Å	10.1 Å	9.5 Å	10.1 Å	
C5	12.3 Å	11.2 Å	10.4 Å	10.9 Å	
C6	13.0 Å	11.8 Å	11.0 Å	11.3 Å	
C7	12.7 Å	11.7 Å	10.9 Å	11.2 Å	
C8	12.9 Å	11.7 Å	10.9 Å	10.9 Å	
C9	11.7 Å	10.4 Å	9.6 Å	9.6 Å	
C10	10.9 Å	9.4 Å	8.5 Å	8.5 Å	
C11	9.6 Å	8.2 Å	7.4 Å	7.5 Å	
C12	8.5 Å	7.3 Å	6.6 Å	6.8 Å	
C13	8.4 Å	7.5 Å	7.1 Å	7.5 Å	
C14	7.2 Å	6.5 Å	6.2 Å	6.9 Å	
C15	6.1 Å	5.2 Å	4.8 Å	5.5 Å	
C16	4.8 Å	4.3 Å	4.2 Å	5.2 Å	
O17	4.5 Å	3.5 Å	3.2 Å	4.4 Å	

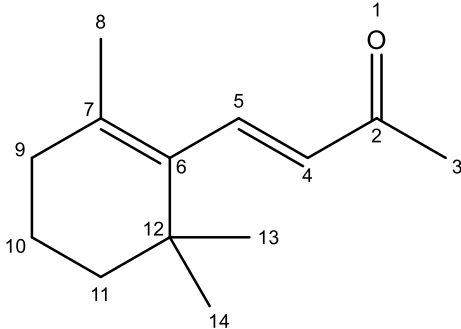
**Table S7.** The distance between the partial carbon atoms on the amino acid residues of OachGOBP2 and each Z3-Hexen-1-ol atom.

OachGOBP2 Z3-Hexen-1-ol atom	THR88-C	ARG132-CZ	Structure
C1	10.3 Å	10.3 Å	
C2	8.8 Å	9.2 Å	
C3	8.3 Å	8.3 Å	
C4	7.1 Å	7.0 Å	
C5	6.0 Å	6.3 Å	
C6	5.3 Å	4.8 Å	
O7	3.9 Å	4.2 Å	

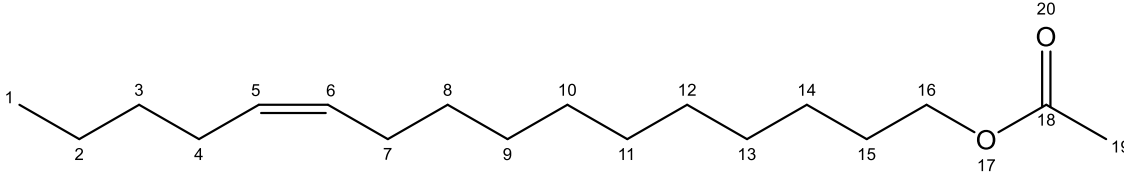
**Table S8.** The distance between the partial carbon atoms on the amino acid residues of OachGOBP2 and each Linalool atom.

OachGOBP2 Linalool atom	THR31-CA	THR31-CB	THR31-CG2	TRP59-CE2	TRP59-CZ2	Structure
C1	4.8 Å	4.7 Å	4.8 Å	4.6 Å	4.3 Å	
C2	4.5 Å	4.8 Å	4.9 Å	6.0 Å	5.8 Å	
C3	4.9 Å	5.6 Å	6.0 Å	6.9 Å	6.4 Å	
C4	6.2 Å	6.9 Å	7.2 Å	8.2 Å	7.7 Å	
C5	7.4 Å	8.1 Å	8.3 Å	8.6 Å	8.0 Å	
C6	7.8 Å	8.2 Å	8.5 Å	8.0 Å	7.2 Å	
C7	8.7 Å	9.4 Å	9.5 Å	10.0 Å	9.4 Å	
C8	6.2 Å	6.0 Å	5.8 Å	4.9 Å	4.6 Å	
C9	7.0 Å	6.9 Å	6.5 Å	6.1 Å	5.9 Å	
C10	5.1 Å	4.9 Å	5.4 Å	4.3 Å	3.7 Å	
O11	4.1 Å	3.7 Å	3.5 Å	3.7 Å	3.9 Å	

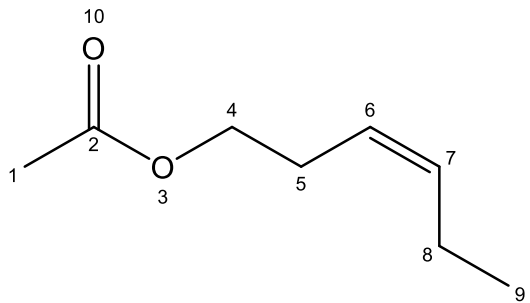
**Table S9.** The distance between the partial carbon atoms on the amino acid residues of OachGOBP2 and each  $\beta$ -Ionone atom.

<b>OachGOBP2 <math>\beta</math>-Ionone atom</b>	<b>SER78-C</b>	<b>Structure</b>
<b>O1</b>	4.1 Å	
<b>C2</b>	5.0 Å	
<b>C3</b>	5.9 Å	
<b>C4</b>	5.7 Å	
<b>C5</b>	5.5 Å	
<b>C6</b>	6.5 Å	
<b>C7</b>	6.8 Å	
<b>C8</b>	6.3 Å	
<b>C9</b>	8.1 Å	
<b>C10</b>	8.9 Å	
<b>C11</b>	9.0 Å	
<b>C12</b>	7.7 Å	
<b>C13</b>	7.4 Å	
<b>C14</b>	8.3 Å	

**Table S10.** The distance between the partial carbon atoms on the amino acid residues of OachGOBP2 and each Z11-16: OAc atom.

OachGOBP2		Structure
Z11-16: OAc atom	SER78-CB	
C1	7.4 Å	
C2	7.7 Å	
C3	8.4 Å	
C4	8.2 Å	
C5	7.8 Å	
C6	8.2 Å	
C7	9.1 Å	
C8	9.5 Å	
C9	10.6 Å	
C10	10.3 Å	
C11	9.9 Å	
C12	8.5 Å	
C13	7.8 Å	
C14	6.3 Å	
C15	6.2 Å	
C16	5.8 Å	
O17	4.5 Å	
C18	4.6 Å	
C19	5.8 Å	
O20	4.1 Å	

**Table S11.** The distance between the partial carbon atoms on the amino acid residues of OachGOBP2 and each Z3-Hexenyl acetate atom.

Z3-Hexenyl acetate atom	OachGOBP2	THR31-CA	THR31-CB	THR31-CG2	Structure
C1		4.4 Å	5.1 Å	5.5 Å	
C2		3.7 Å	4.0 Å	4.5 Å	
O3		4.1 Å	4.1 Å	4.1 Å	
C4		5.3 Å	4.9 Å	4.9 Å	
C5		6.5 Å	6.3 Å	6.2 Å	
C6		6.8 Å	6.6 Å	6.2 Å	
C7		7.6 Å	7.7 Å	7.2 Å	
C8		8.4 Å	8.6 Å	8.3 Å	
C9		8.0 Å	8.5 Å	8.4 Å	
O10		3.5 Å	3.7 Å	4.6 Å	