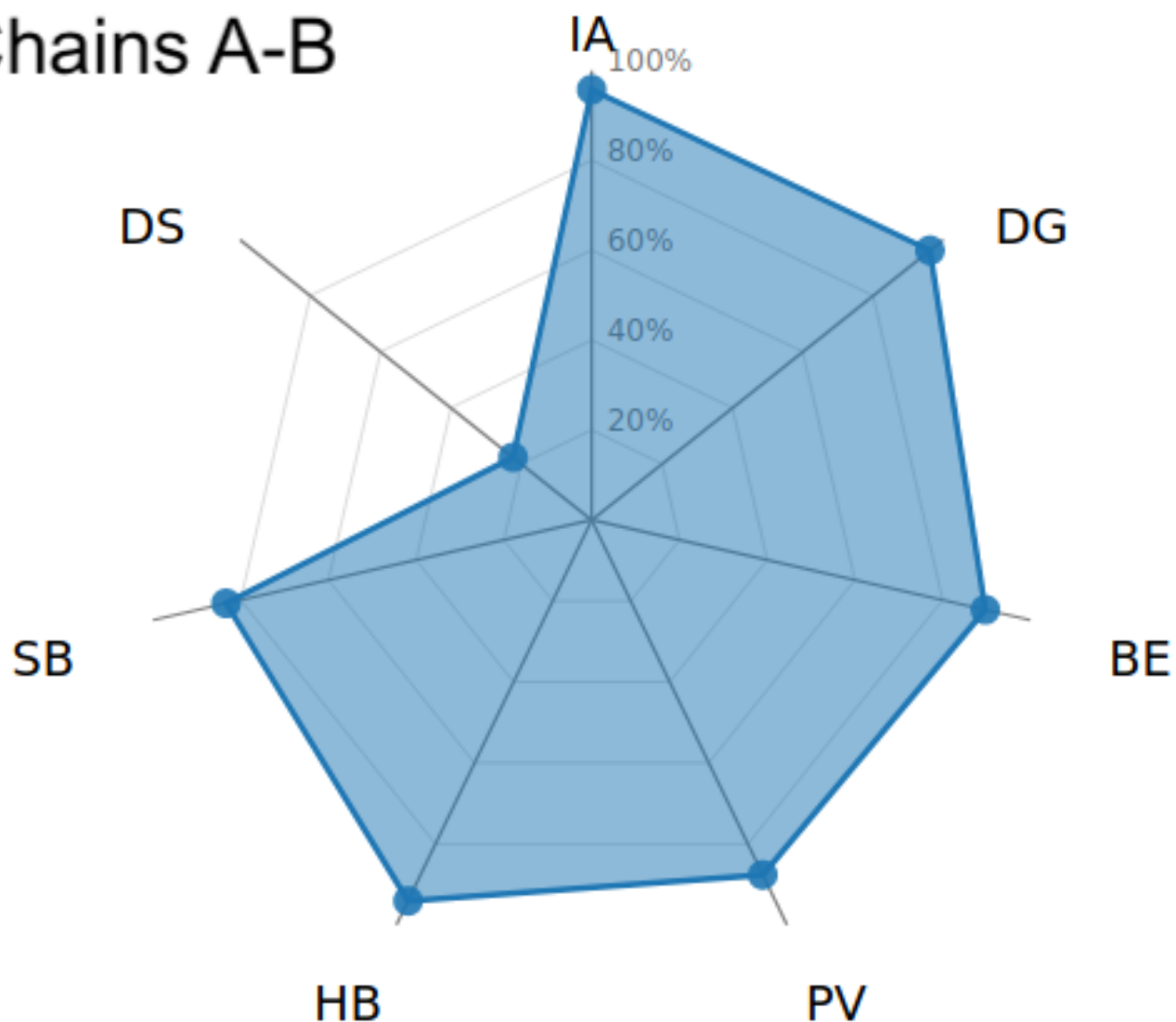


## *Interaction radar*

### Chains A-B

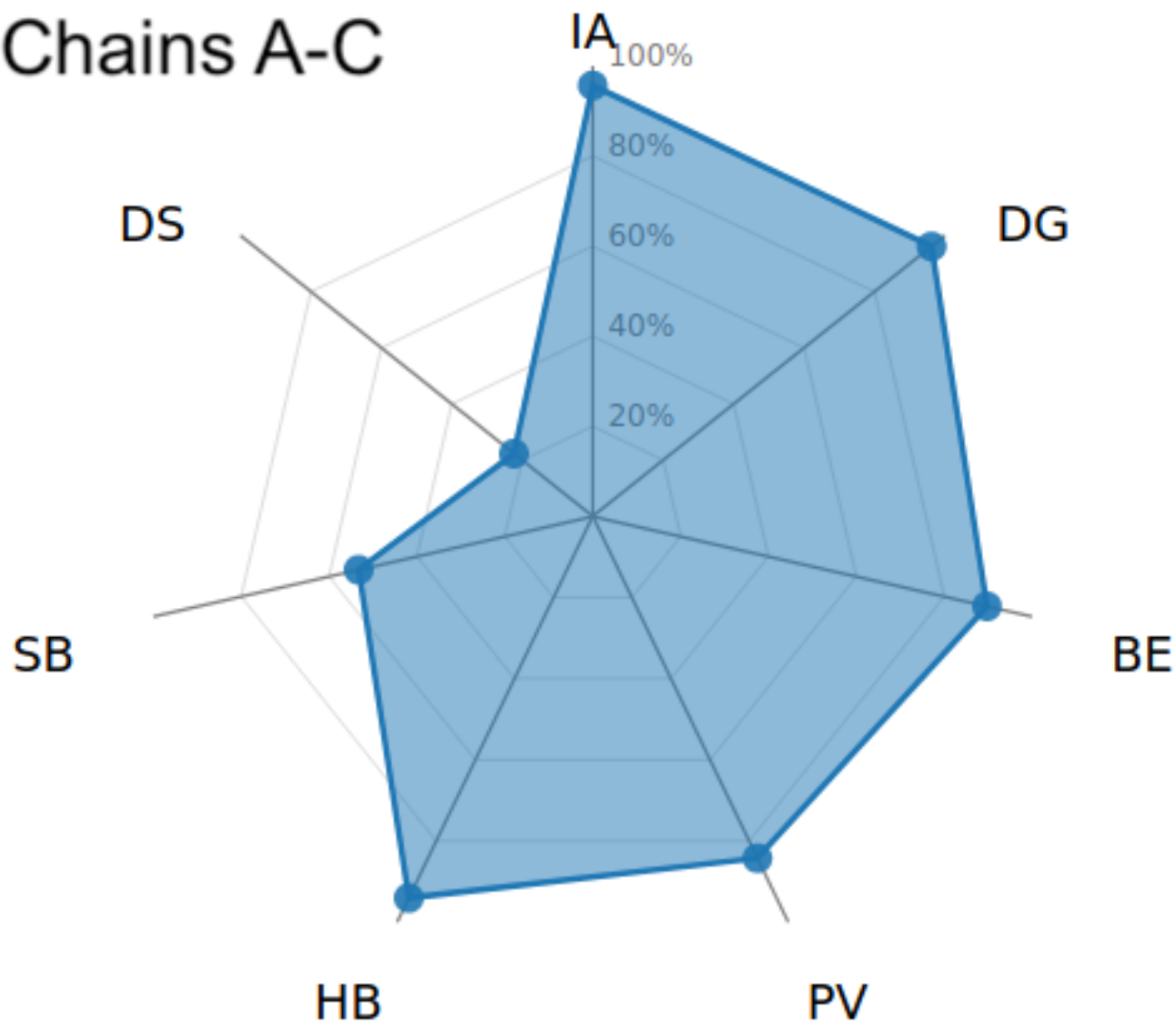


## *Interface parameters*

<b>IA</b>	: Interface area, Å <sup>2</sup>	6968
<b>DG</b>	: Solvation Energy, kcal/mol	-89.34
<b>BE</b>	: Total Binding Energy, kcal/mol	-106.5
<b>PV</b>	: Hydrophobic P-value	0.01455
<b>HB</b>	: Number of Hydrogen Bonds	26
<b>SB</b>	: Number of Salt Bridges	15
<b>DS</b>	: Number of Disulphide Bonds	0

## *Interaction radar*

### Chains A-C

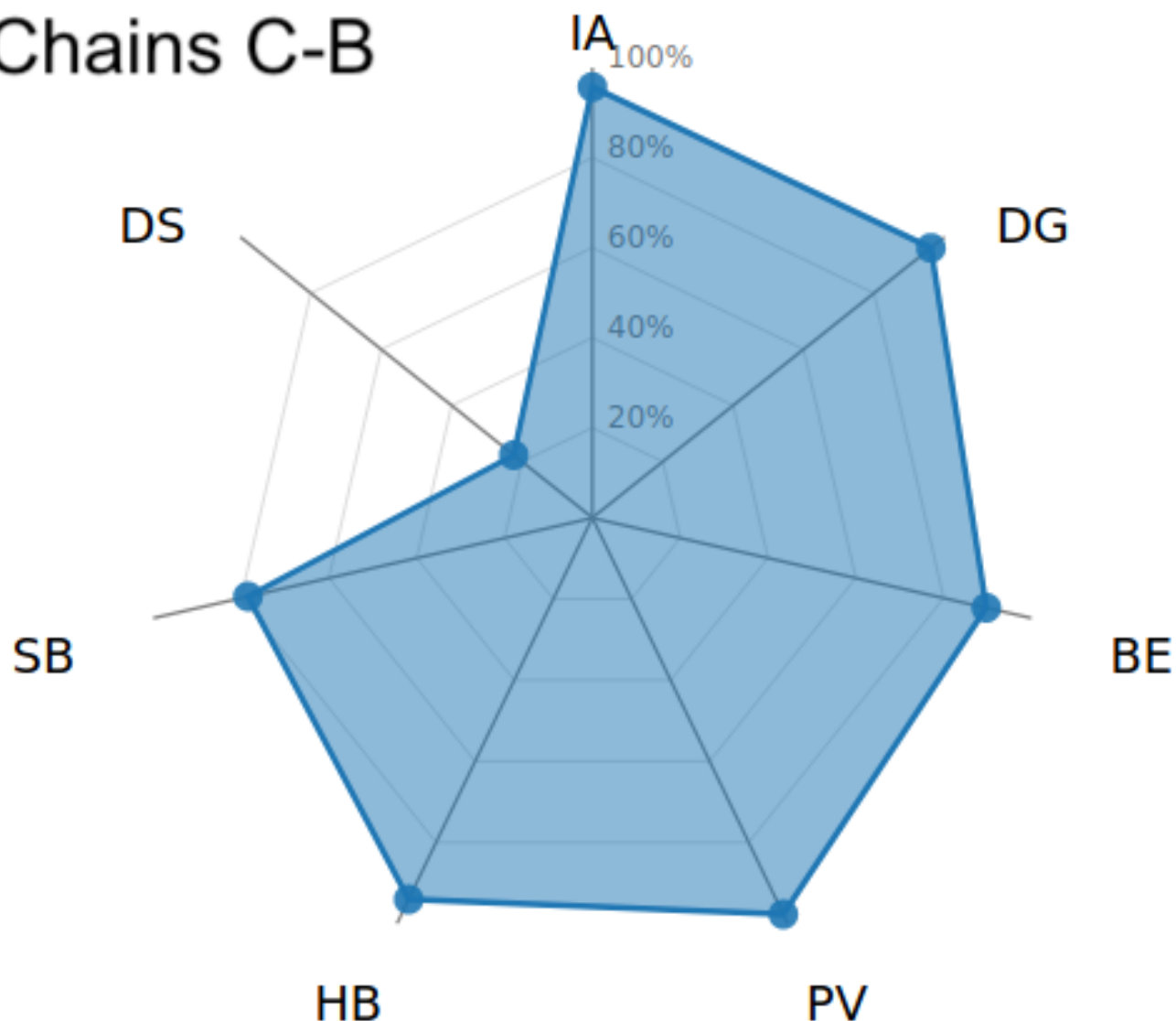


## *Interface parameters*

<b>IA</b>	: Interface area, Å <sup>2</sup>	6876
<b>DG</b>	: Solvation Energy, kcal/mol	-81.38
<b>BE</b>	: Total Binding Energy, kcal/mol	-97.75
<b>PV</b>	: Hydrophobic P-value	0.03473
<b>HB</b>	: Number of Hydrogen Bonds	31
<b>SB</b>	: Number of Salt Bridges	7
<b>DS</b>	: Number of Disulphide Bonds	0

## Interaction radar

### Chains C-B



### Interface parameters

<b>IA</b>	: Interface area, Å <sup>2</sup>	6175
<b>DG</b>	: Solvation Energy, kcal/mol	-90.7
<b>BE</b>	: Total Binding Energy, kcal/mol	-105.2
<b>PV</b>	: Hydrophobic P-value	0.002389
<b>HB</b>	: Number of Hydrogen Bonds	21
<b>SB</b>	: Number of Salt Bridges	14
<b>DS</b>	: Number of Disulphide Bonds	0