

Working with the SAME command

There are two ways to put the SAME instruction into your model in Olex2

1. Select reference group in a particular order, then select the dependent group(s) in the same order and type:

```
same n
```

where n is the number of groups (would be 2 if there is only one dependent group).

2. If the molecules can be matched through using 'match' command, you can select an atom in the reference molecule and an atom in dependent molecule and type:

```
same 2
```

In this case also note if the inversion was used in the 'match' command.