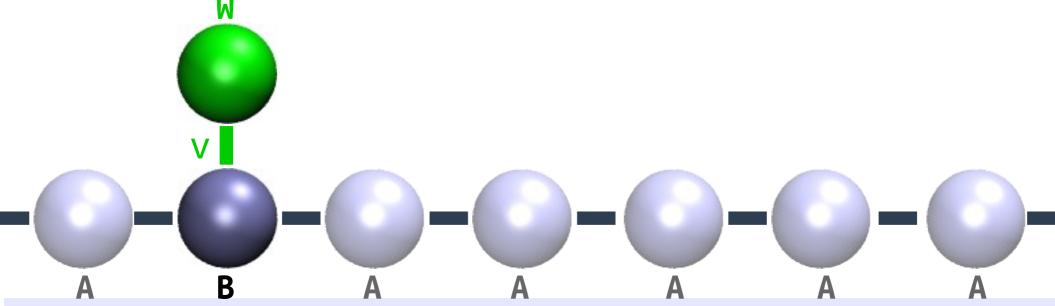


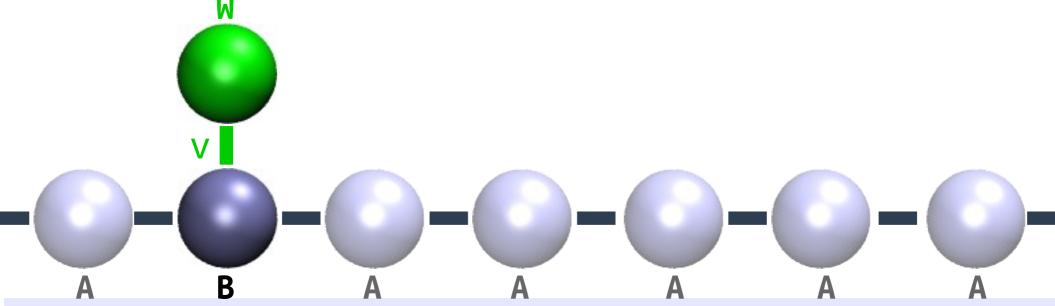
```
if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt

fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V

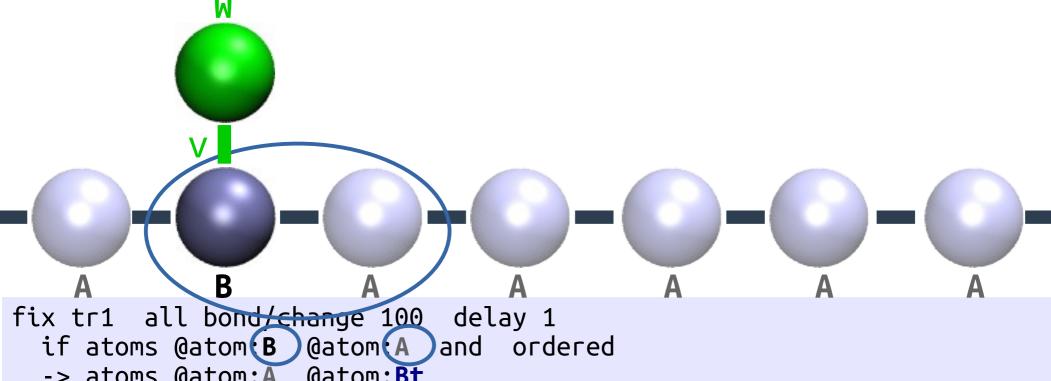
fix tr3 all bond/modify 100 delay 3
  if atoms @atom:Wt @atom:A
  -> atoms @atom:W @atom:A and bond BREAK
```



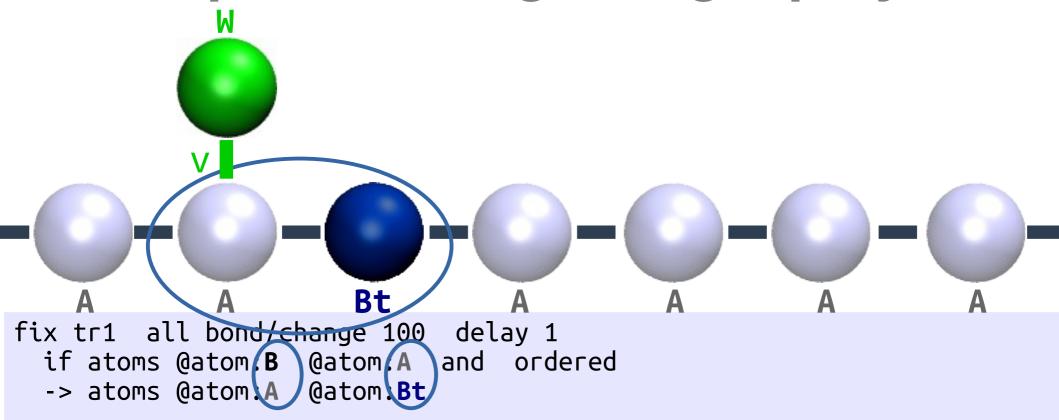
```
fix tr1 all bond/change 100 delay 1
  if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt
```

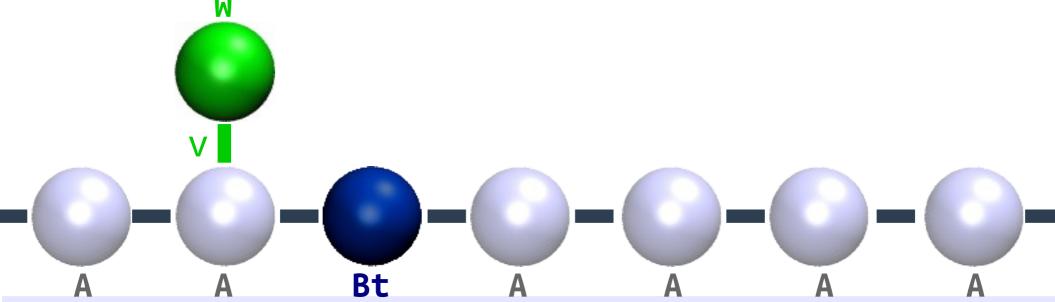


```
fix tr1 all bond/change 100 delay 1
  if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt
```

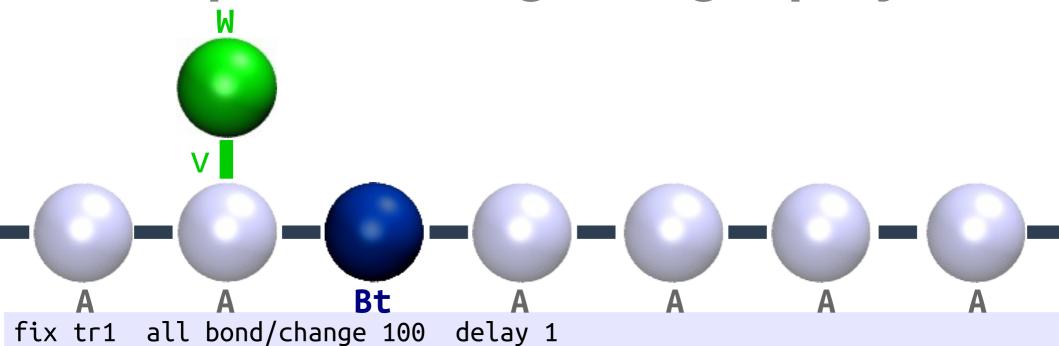


-> atoms @atom:A @atom:Bt



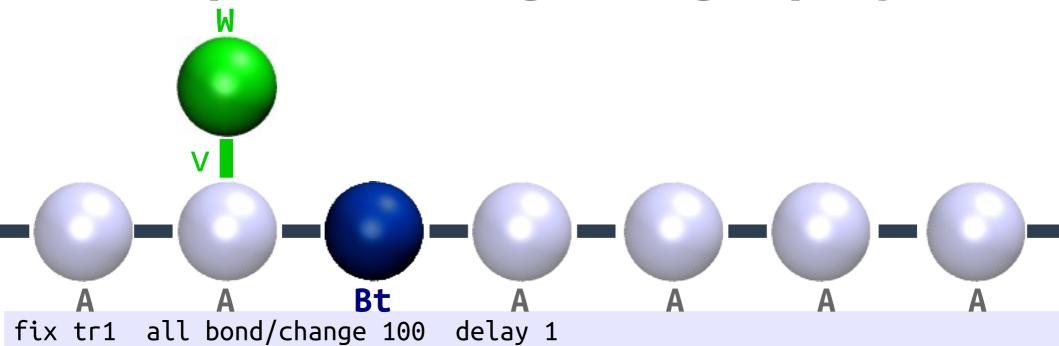


```
fix tr1 all bond/change 100 delay 1
  if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt
```



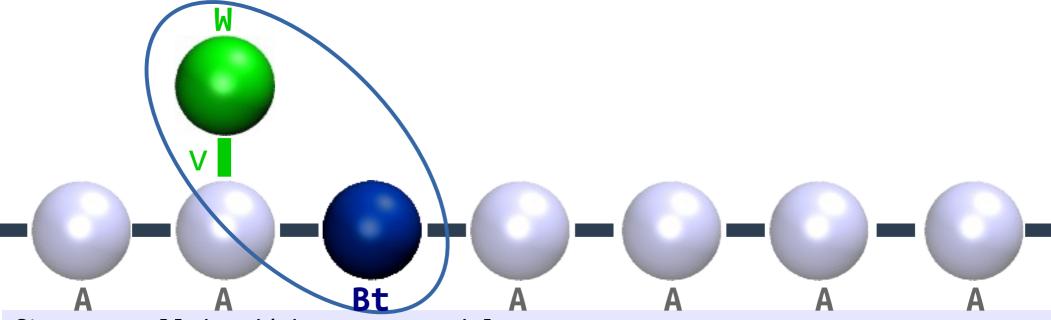
```
if atoms @atom:B @atom:A and ordered
-> atoms @atom:A @atom:Bt
```

```
fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V
```



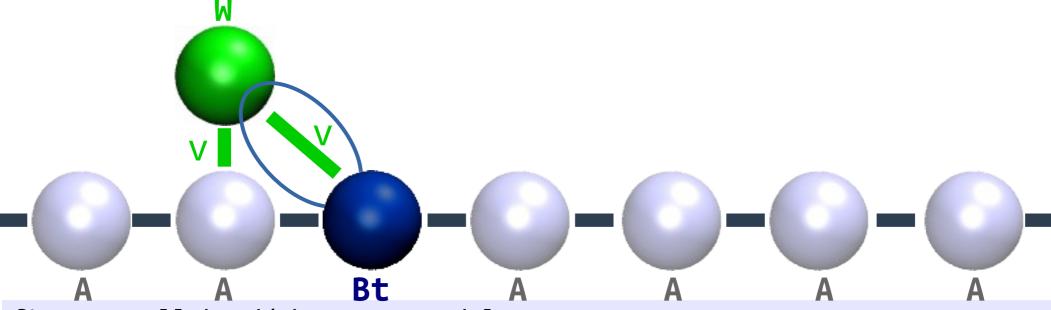
```
-> atoms @atom:A @atom:Bt
fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V
```

if atoms @atom: A and ordered



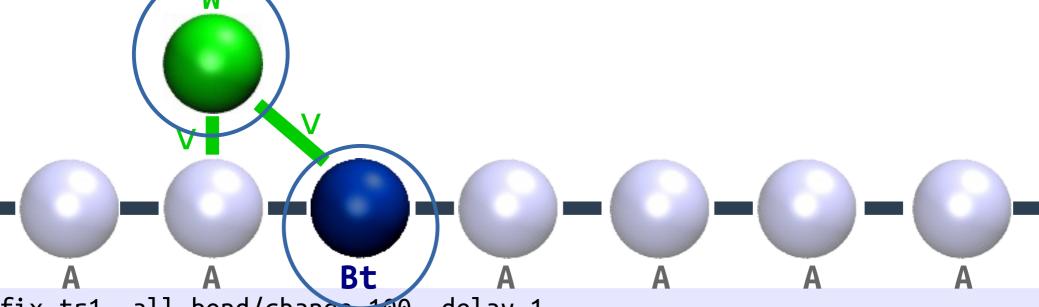
```
fix tr1 all bond/change 100 delay 1
  if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt
```

```
fix tr2 all bond/new 100 delay 2
  if atoms @atom(W) @atom(Bt) and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V
```



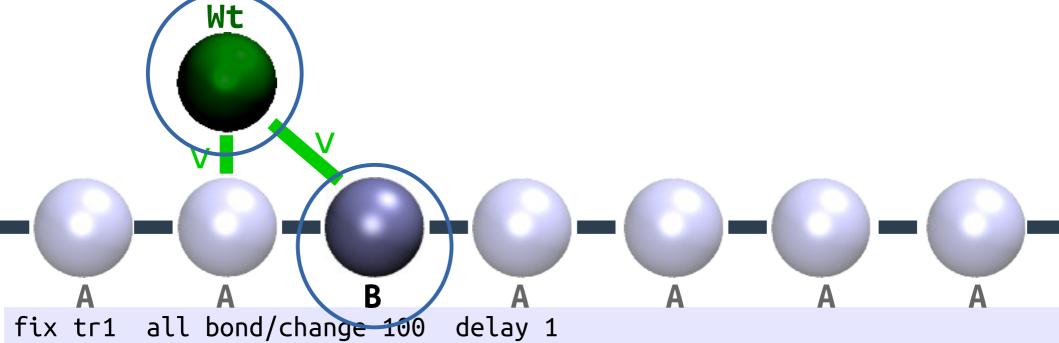
```
fix tr1 all bond/change 100 delay 1
  if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt
```

```
fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V
```



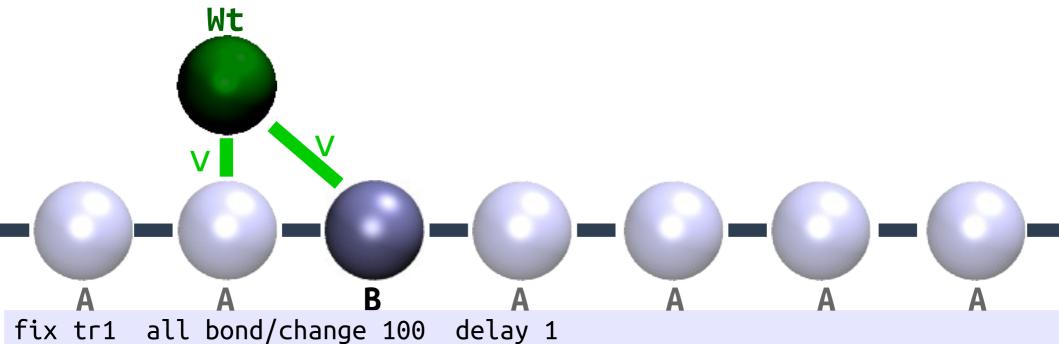
```
fix tr1 all bond/change 100 delay 1
  if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt
```

```
fix tr2 all bond/new 100 delay 2
if atoms @atom W @atom B and distance <= 1.5
-> atoms @atom Wt @atom B and bond @bond:V
```



```
if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt
```

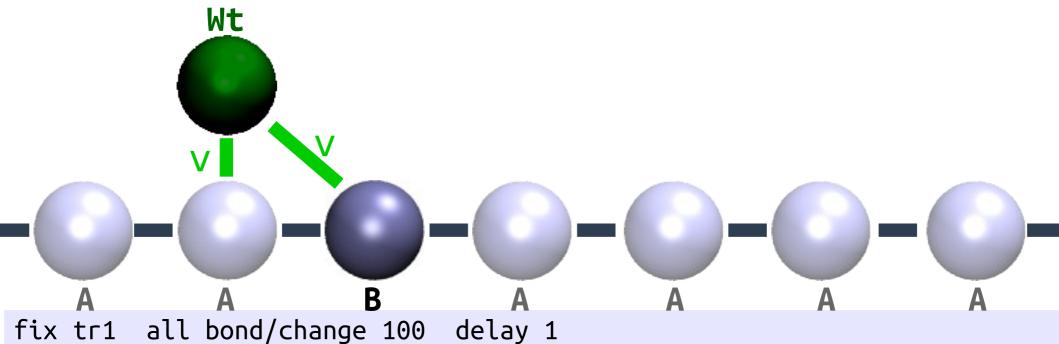
```
fix tr2 all bond/new 100 delay 2
if atoms @atom W @atom B and distance <= 1.5
-> atoms @atom Wt @atom B and bond @bond:V
```



```
if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt

fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5</pre>
```

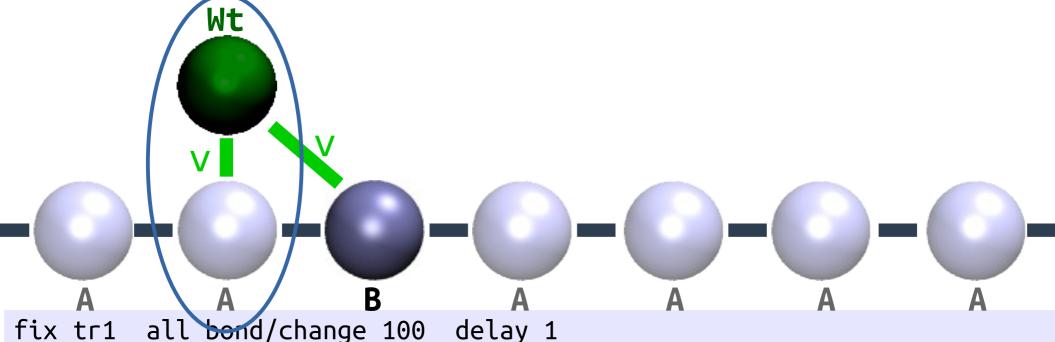
-> atoms @atom:Wt @atom:B and bond @bond:V



```
if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt

fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V

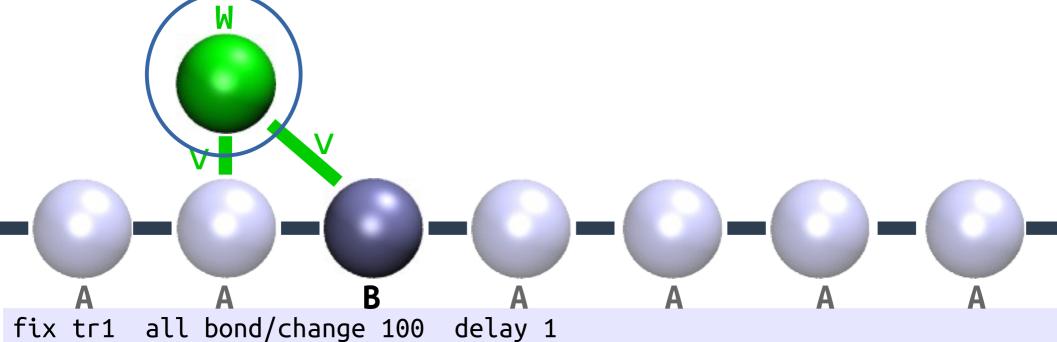
fix tr3 all bond/modify 100 delay 3
  if atoms @atom:Wt @atom:A
  -> atoms @atom:W @atom:A and bond BREAK
```



```
if atoms @atom:B @atom:A and ordered
-> atoms @atom:A @atom:Bt
```

```
fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V
```

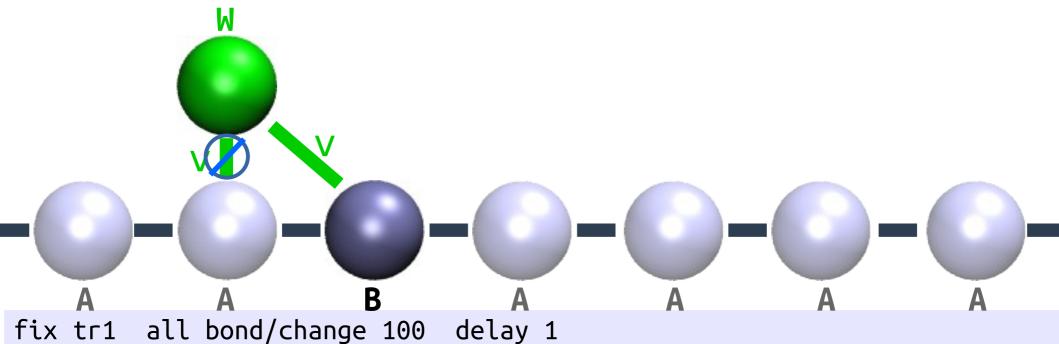
```
fix tr3 all bond/modify 100 delay 3
  if atoms @atom Wt @atom A
  -> atoms @atom:W @atom:A and bond BREAK
```



```
if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt

fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V

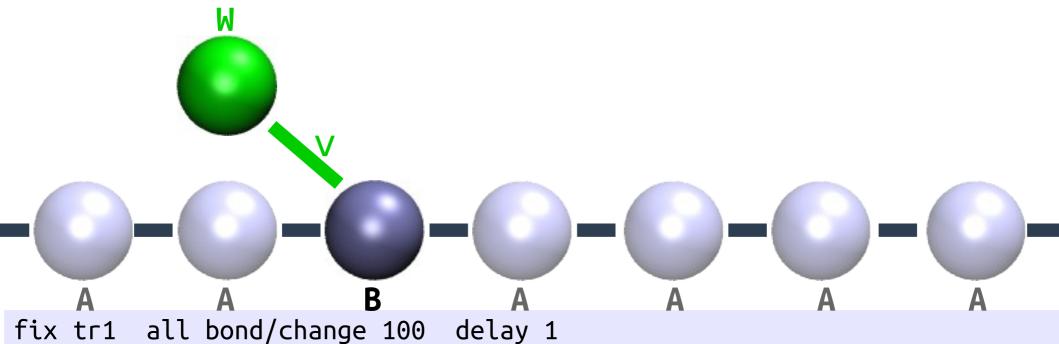
fix tr3 all bond/modify 100 delay 3
  if atoms @atom:Wt @atom:A
  -> atoms @atom:W @atom:A and bond BREAK
```



```
if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt

fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V

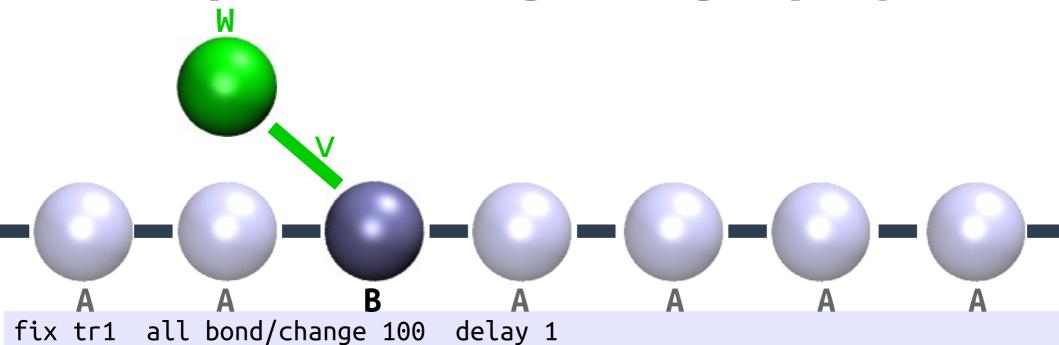
fix tr3 all bond/modify 100 delay 3
  if atoms @atom:Wt @atom:A
  -> atoms @atom:W @atom:A and bond BREAK
```



```
if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt

fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V

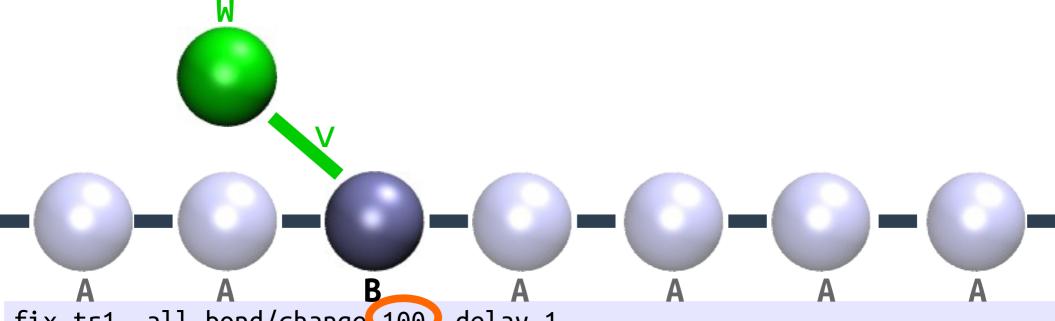
fix tr3 all bond/modify 100 delay 3
  if atoms @atom:Wt @atom:A
  -> atoms @atom:W @atom:A and bond BREAK
```



```
if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt

fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V

fix tr3 all bond/modify 100 delay 3
  if atoms @atom:Wt @atom:A
  -> atoms @atom:W @atom:A and bond BREAK
```

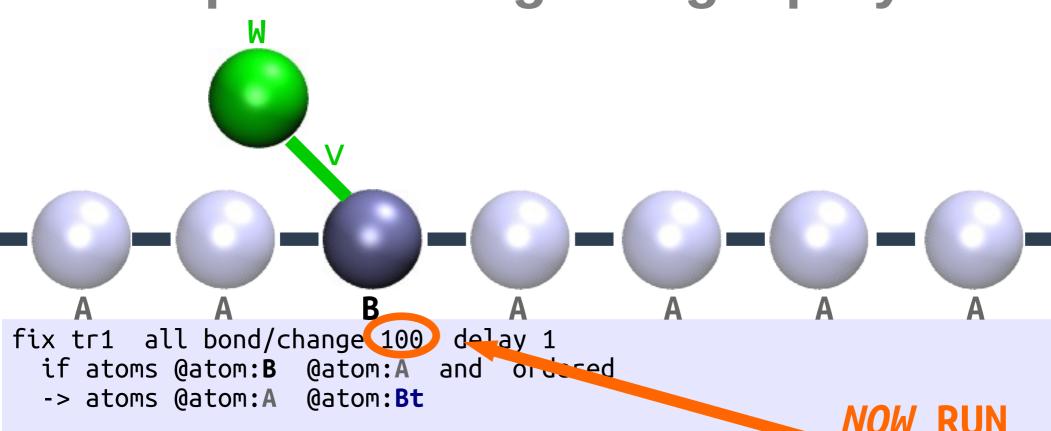


```
fix tr1 all bond/change 100 delay 1
  if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt
```

```
fix tr2 all bond/new 100 delay 2
if atoms @atom:W @atom:Bt and distance <= 1.5
-> atoms @atom:Wt @atom:B and bond @bond.y
```

**NOW** RUN MOLECULAR DYNAMICS

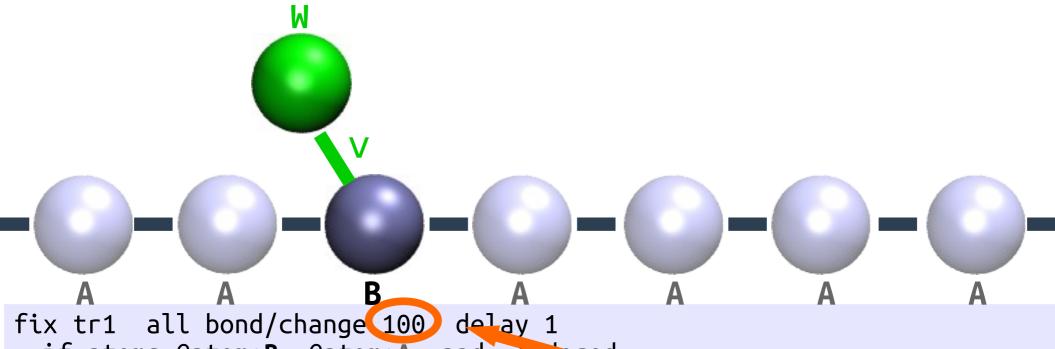
```
fix tr3 all bond/modify 100 detay 3
  if atoms @atom:Wt @atom:A
  -> atoms @atom:W  @atom:A   and  bond BREAK
```



```
fix tr2 all bond/new 100 delay 2
if atoms @atom:W @atom:Bt and distance <= 1.5
-> atoms @atom:Wt @atom:B and bond @bond.y
```

**NOW** RUN MOLECULAR DYNAMICS

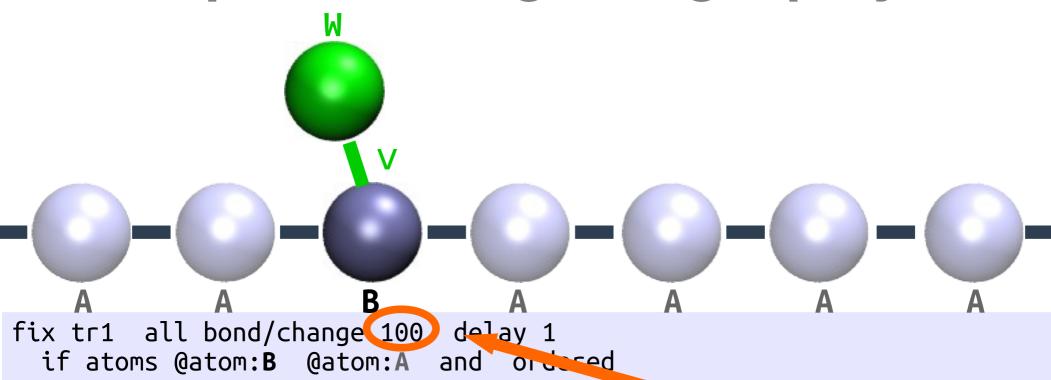
```
fix tr3 all bond/modify 100 detay 3
  if atoms @atom:Wt @atom:A
  -> atoms @atom:W  @atom:A   and  bond BREAK
```



```
fix tr1 all bond/change 100 delay 1
  if atoms @atom:B @atom:A and ordered
  -> atoms @atom:A @atom:Bt
```

```
fix tr2 all bond/new 100 delay 2
if atoms @atom:W @atom:Bt and distance <= 1.5
-> atoms @atom:Wt @atom:B and bond @bond.y
```

 NOW RUN
MOLECULAR
DYNAMICS

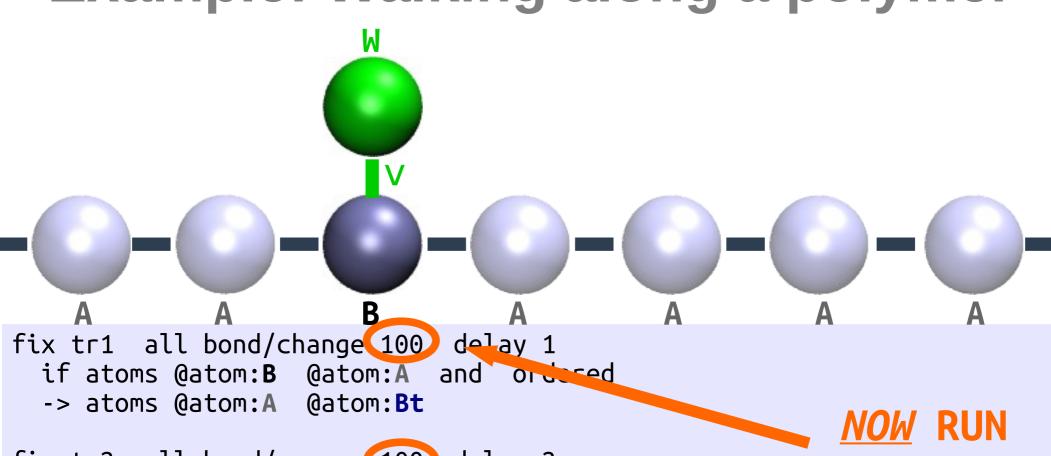


```
-> atoms @atom:A @atom:Bt
```

```
fix tr2 all bond/new
                      100 delay 2
 if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:
```

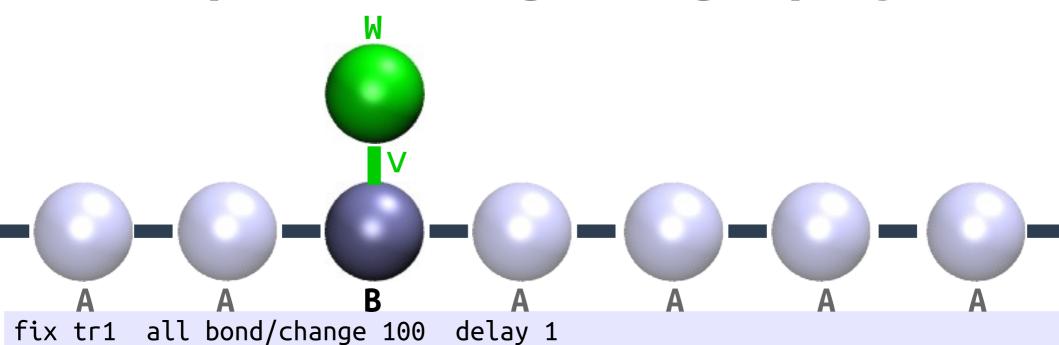
*NOW* RUN **DYNAMICS** 

```
fix tr3 all bond/modify 100 detay 3
 if atoms Qatom: Wt Qatom: A
  -> atoms @atom:W @atom:A and bond BREAK
```



```
fix tr2 all bond/new 100 delay 2
if atoms @atom:W @atom:Bt and distance <= 1.5
-> atoms @atom:Wt @atom:B and bond @bond:V
```

**NOW** RUN MOLECULAR DYNAMICS

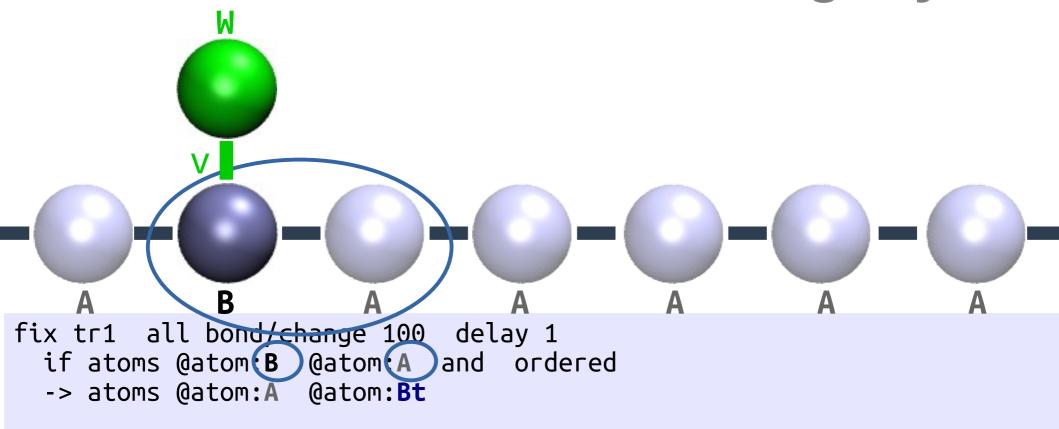


```
if atoms @atom:B @atom:A and ordered
-> atoms @atom:A @atom:Bt

fix tr2 all bond/new 100 delay 2
  if atoms @atom:W @atom:Bt and distance <= 1.5
  -> atoms @atom:Wt @atom:B and bond @bond:V

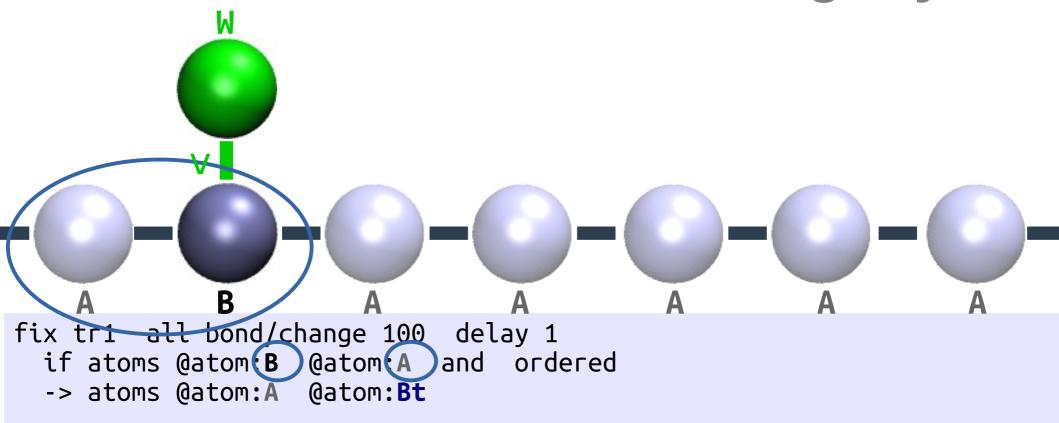
fix tr3 all bond/modify 100 delay 3
  if atoms @atom:Wt @atom:A
  -> atoms @atom:W @atom:A and bond BREAK
```

#### Problem: direction ambiguity



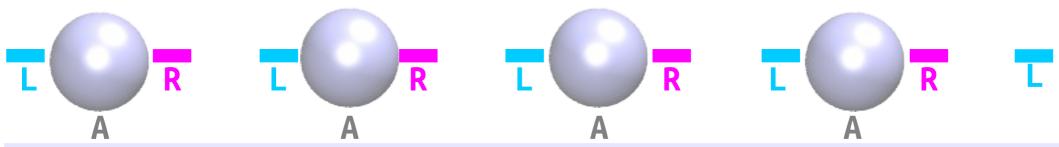
"Walker" can move right or left...

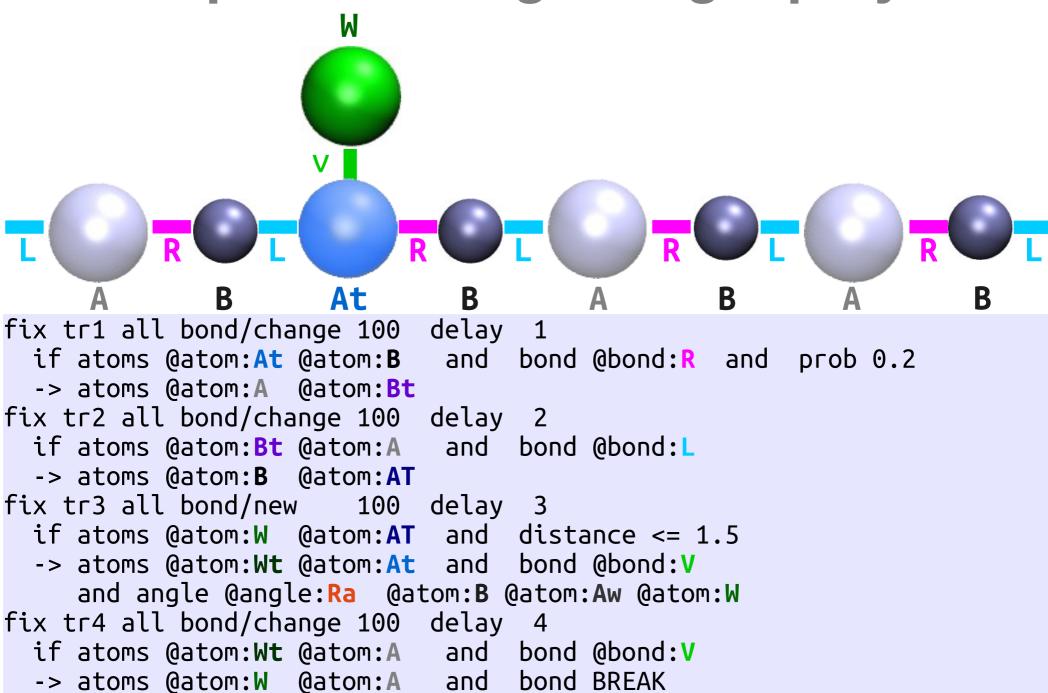
#### Problem: direction ambiguity

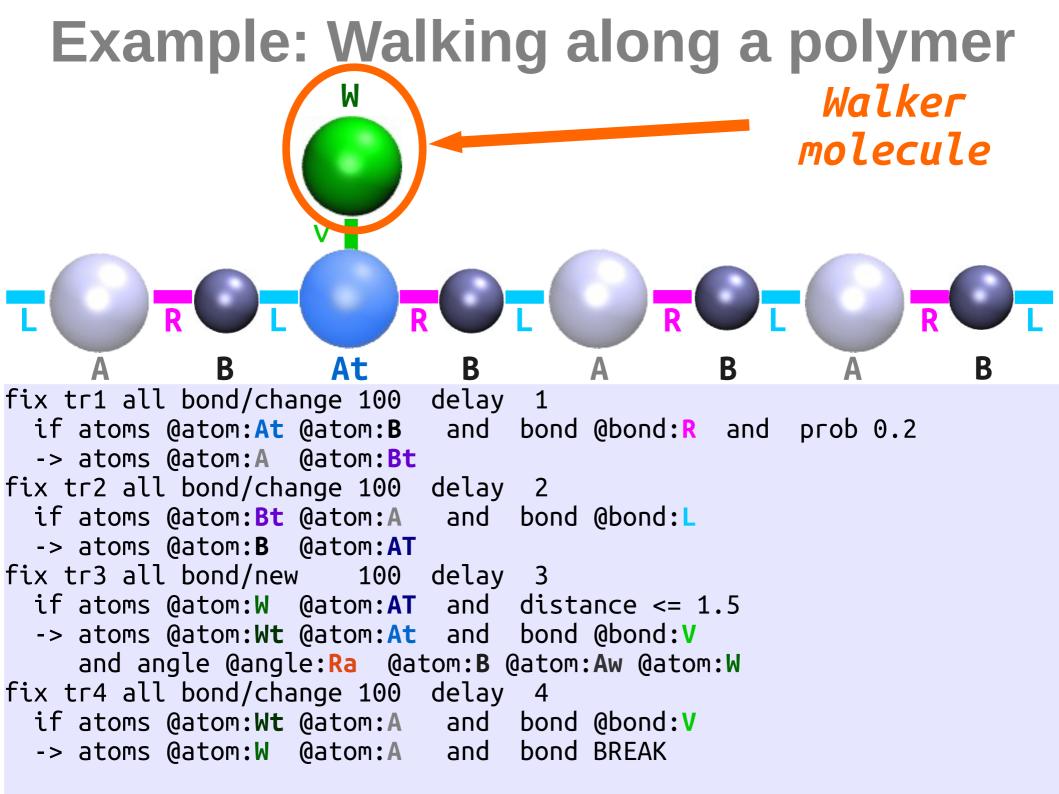


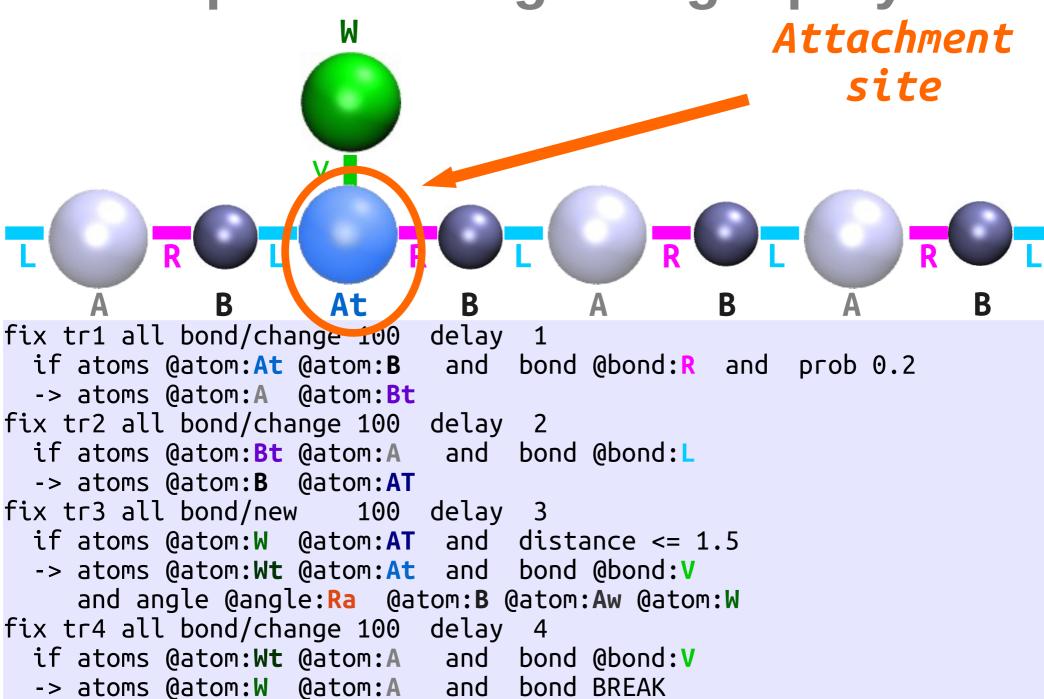
"Walker" can move right or left...

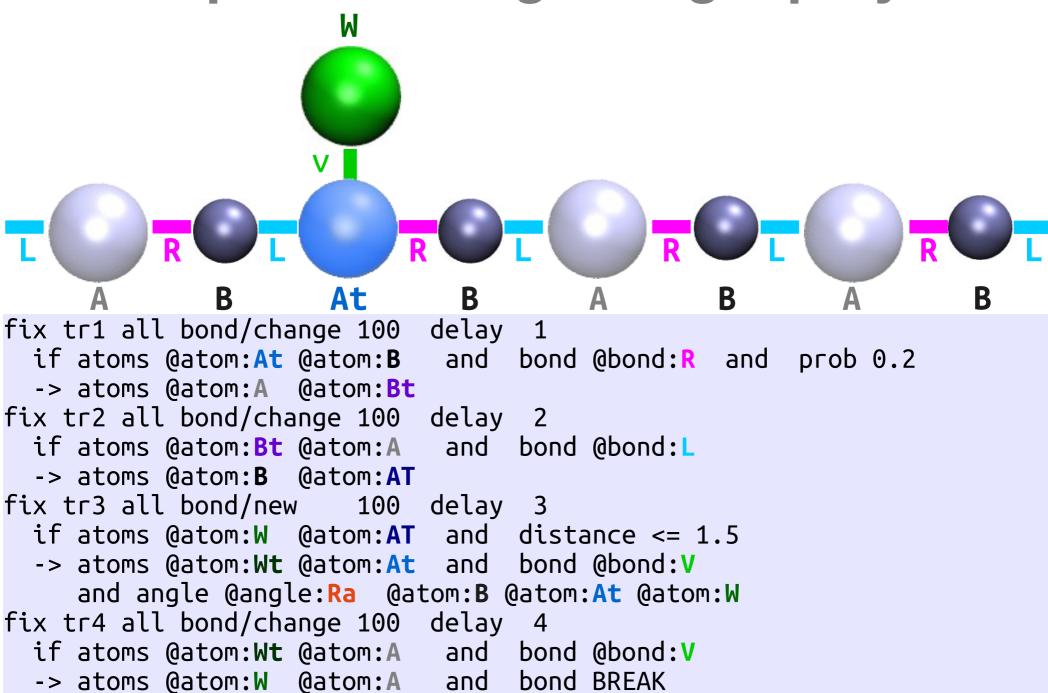
# Solution Strategy: Break symmetry One way to do this is to use different bond types (L and R)

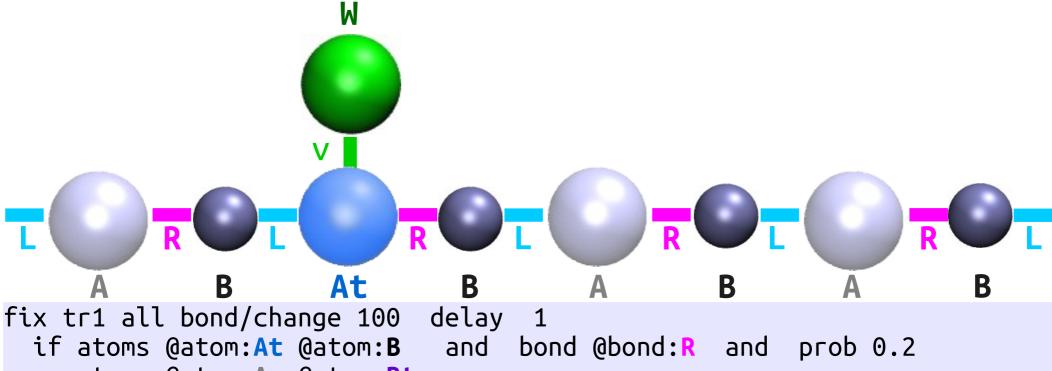




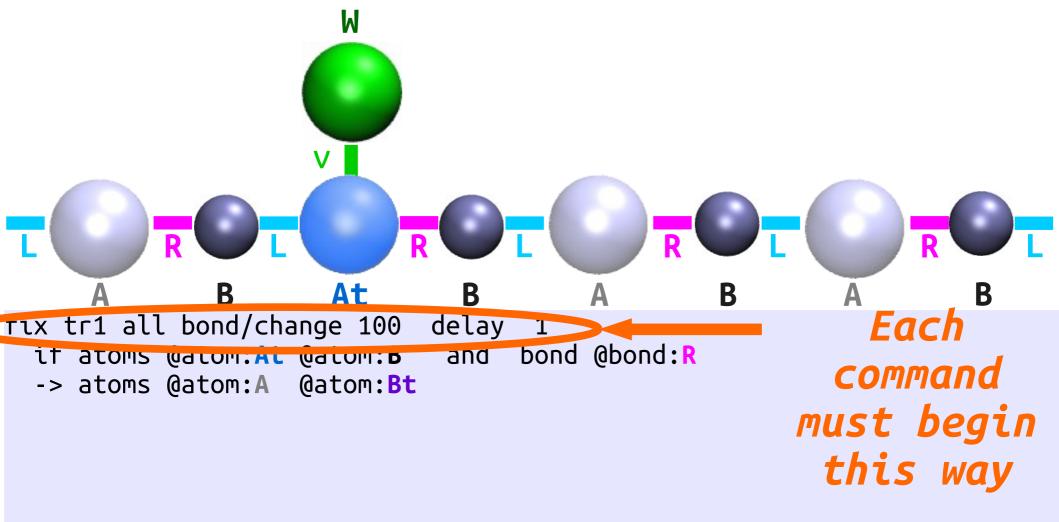




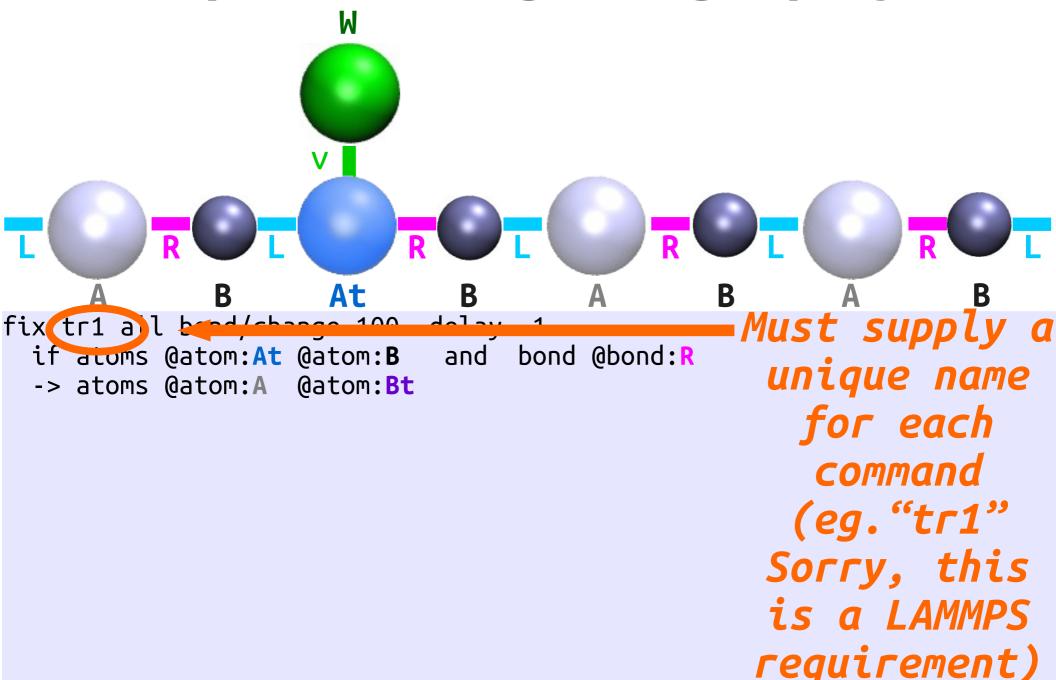


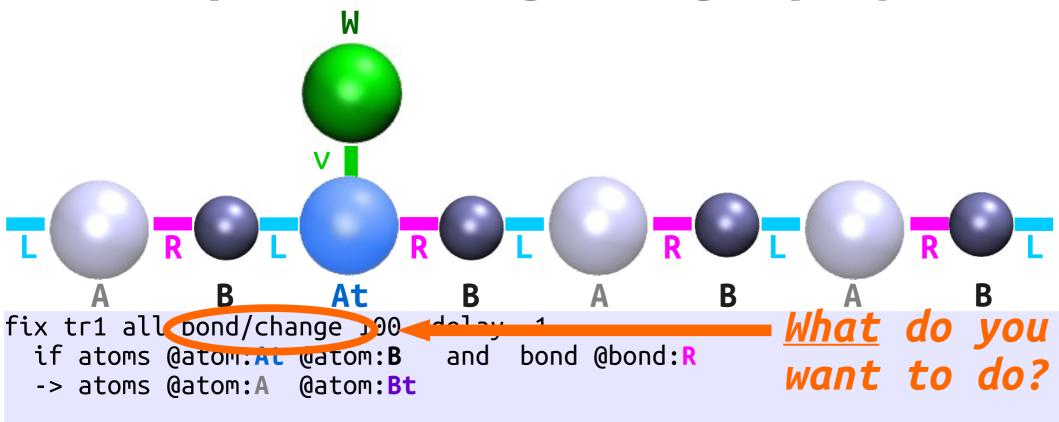


-> atoms @atom:A @atom:Bt

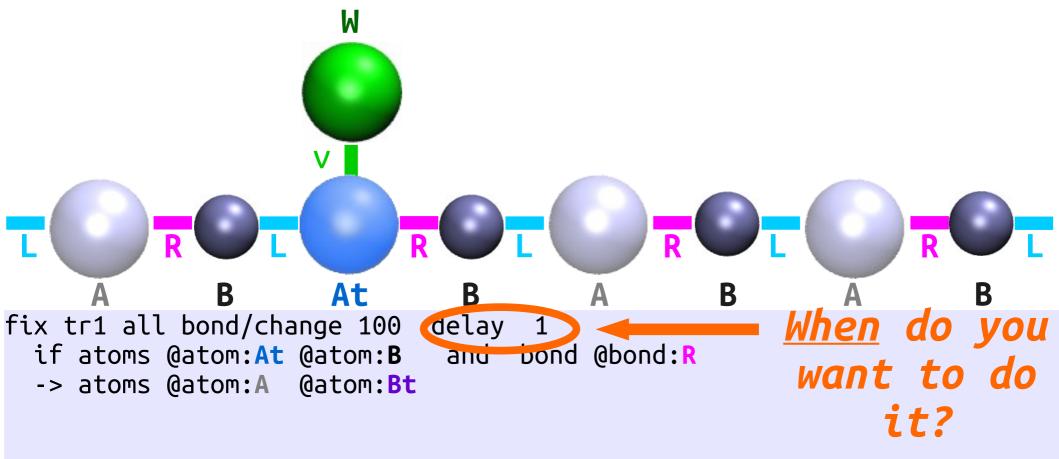


(blame LAMMPS)

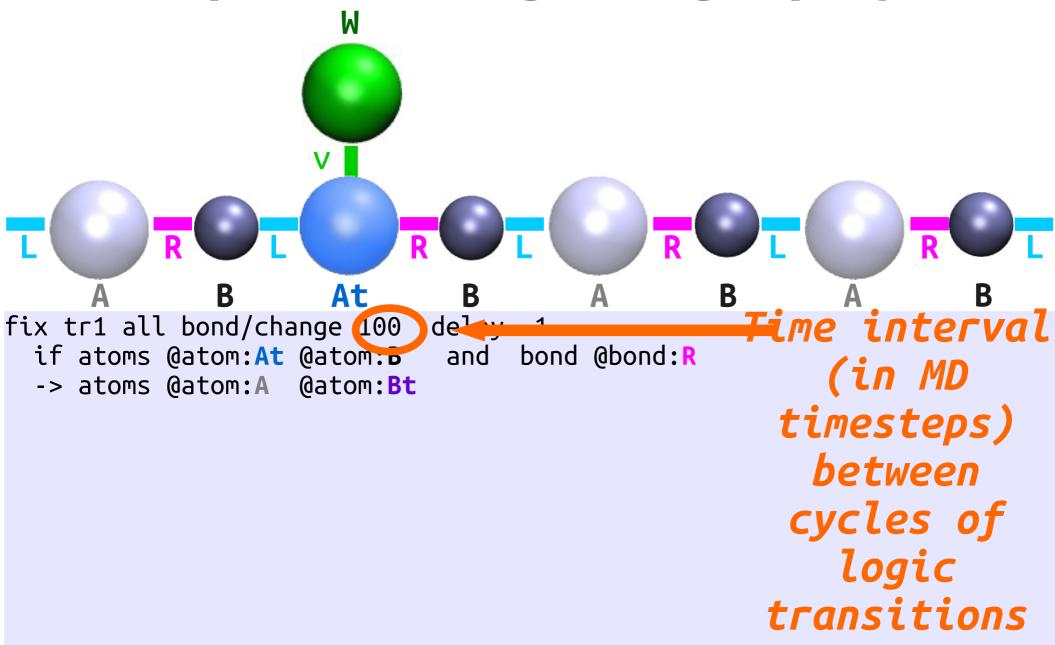


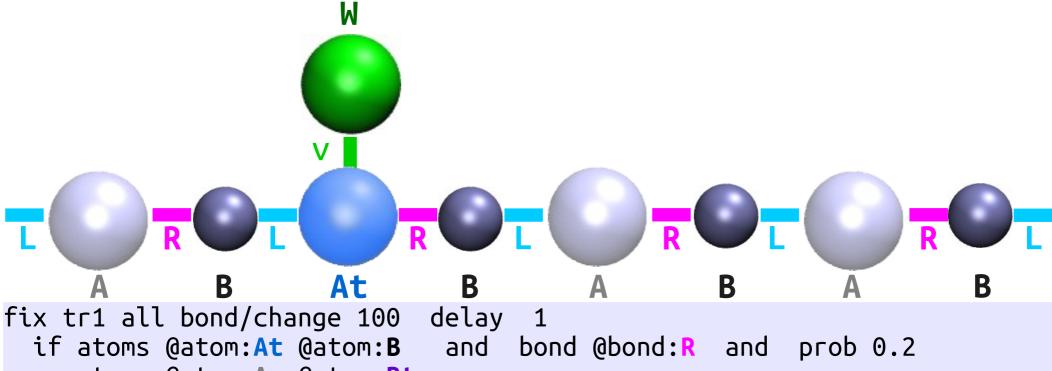


(modify an
 existing
 bond)

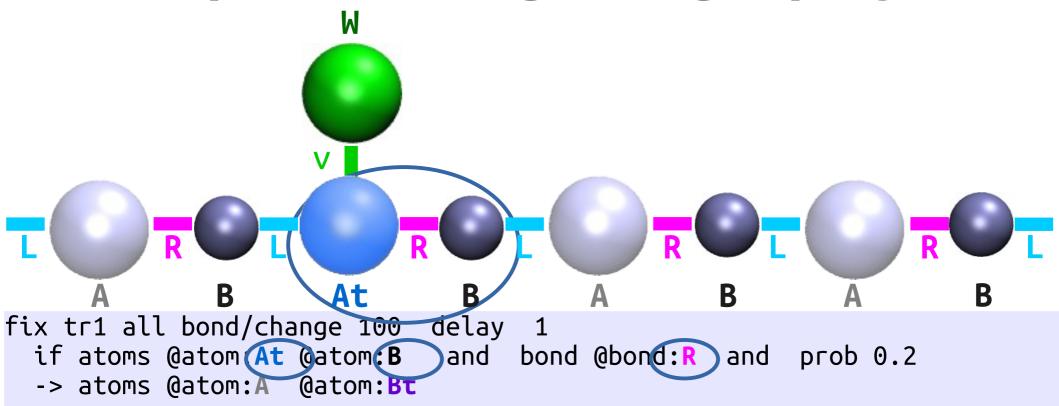


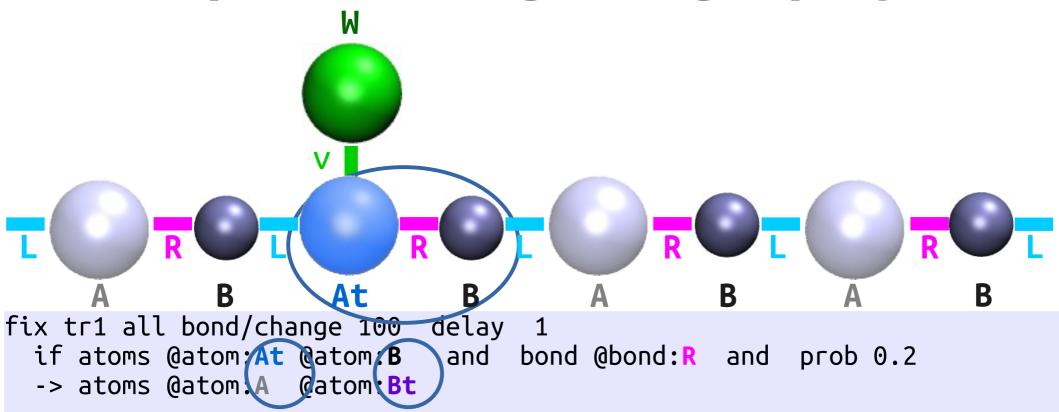
(similar to a "line number")

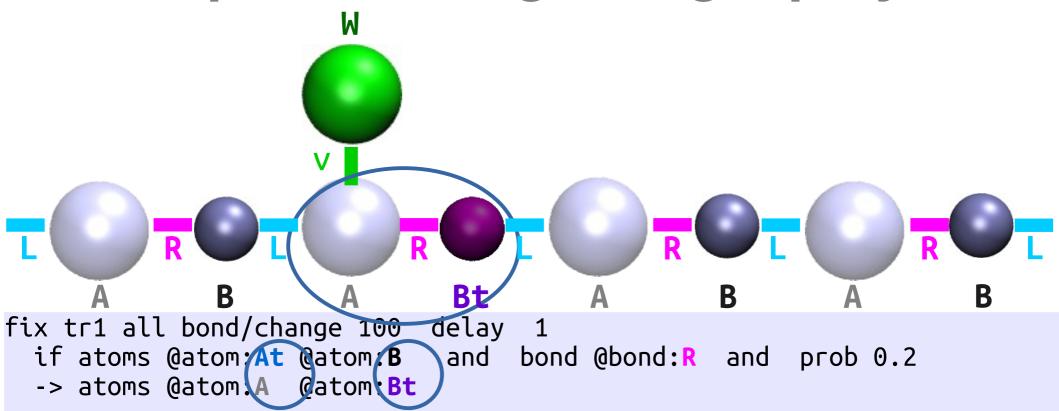


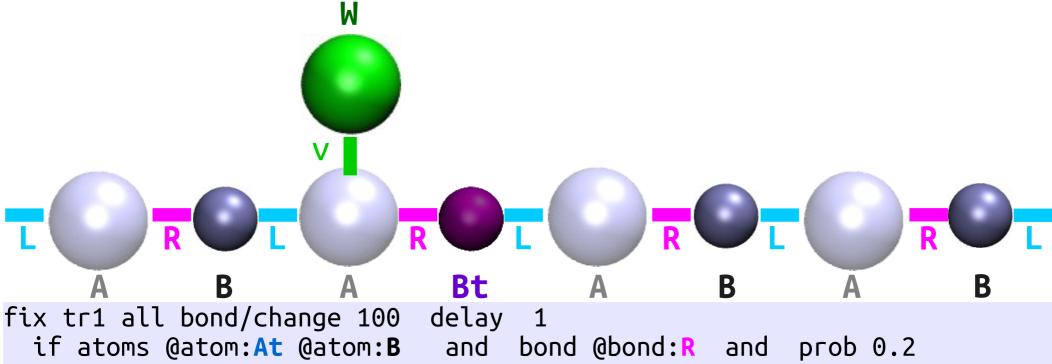


-> atoms @atom:A @atom:Bt

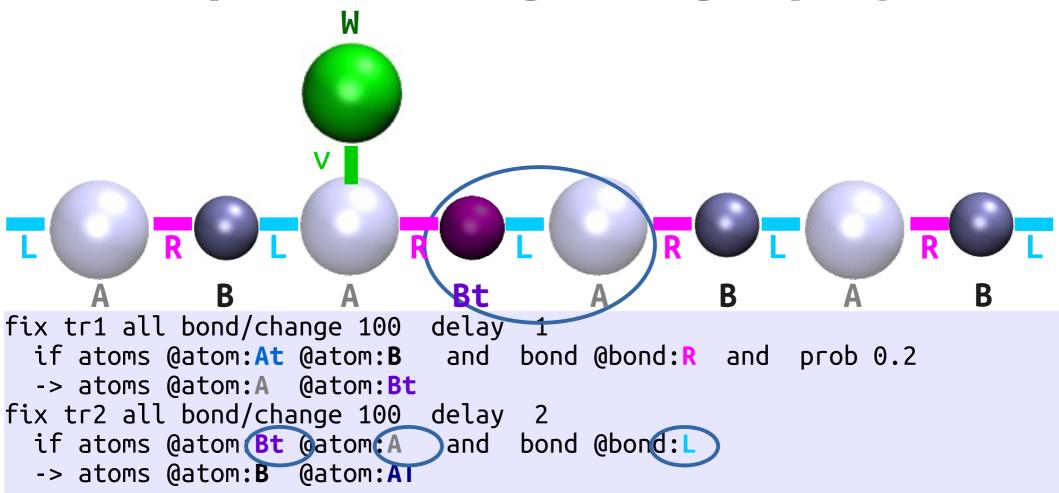


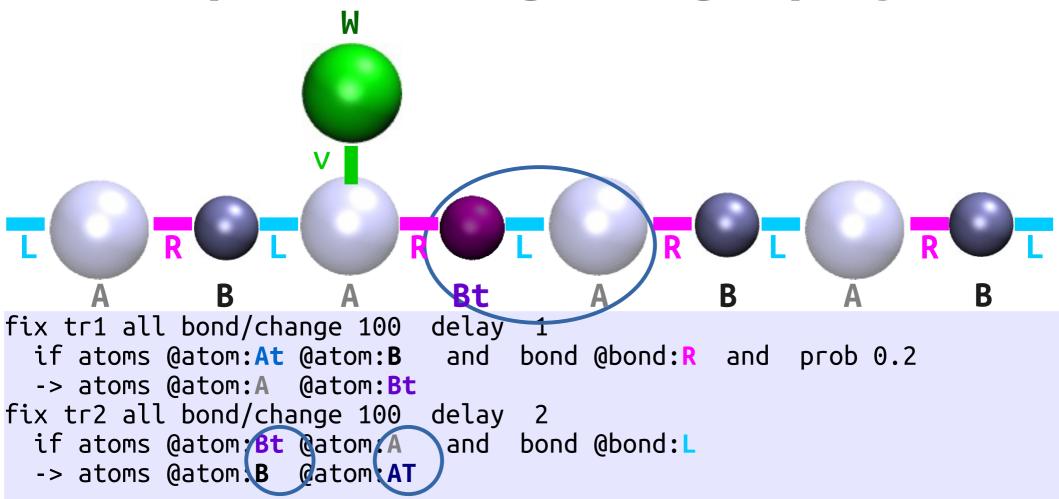


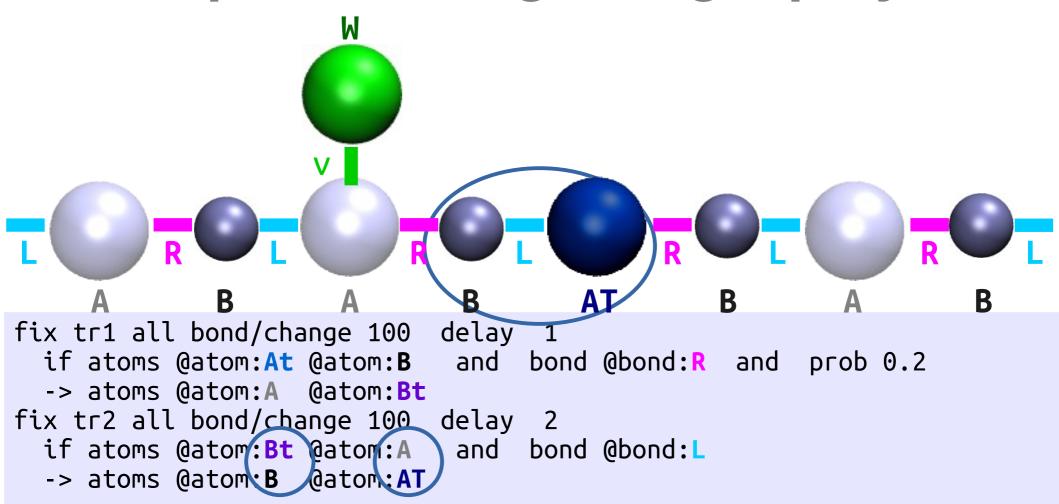


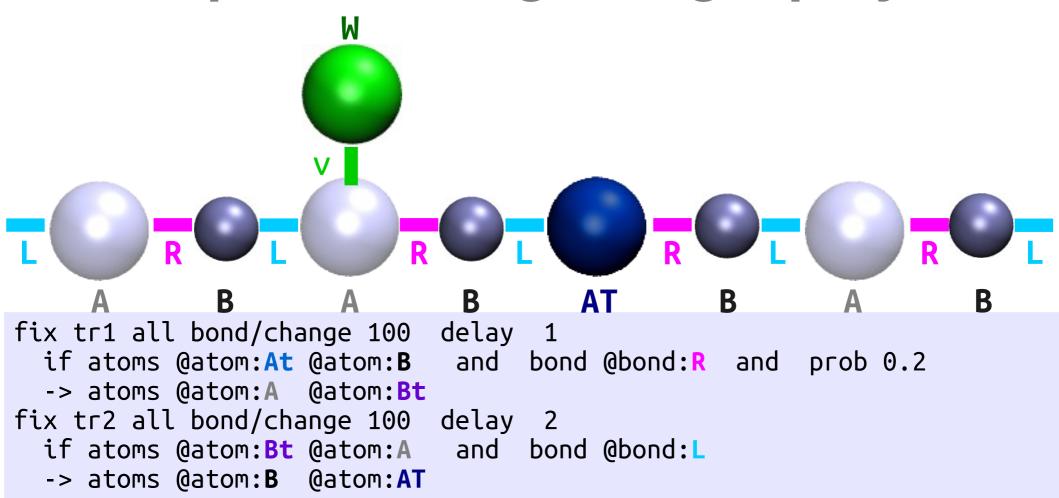


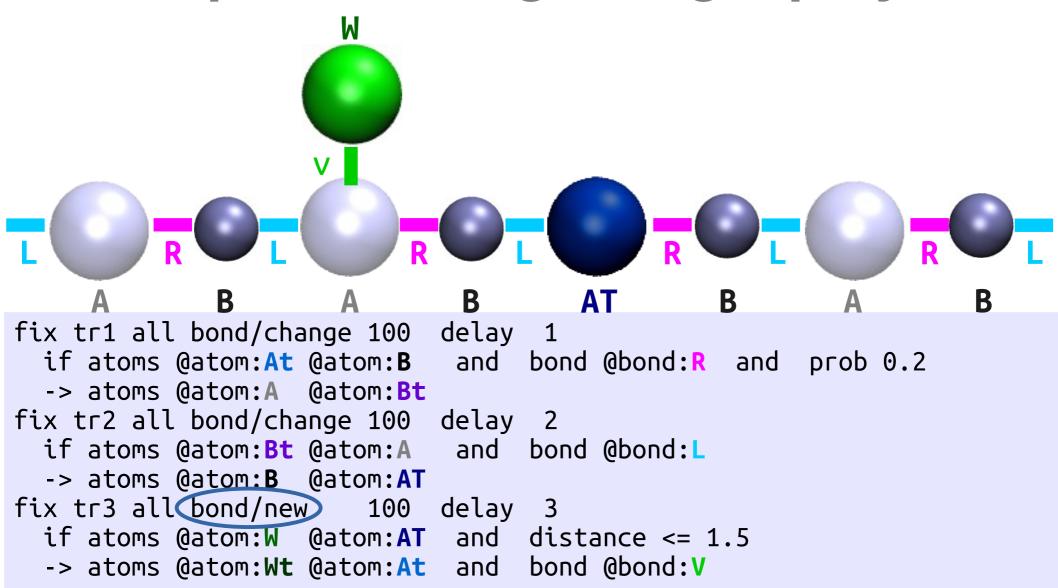
-> atoms @atom:A @atom:Bt

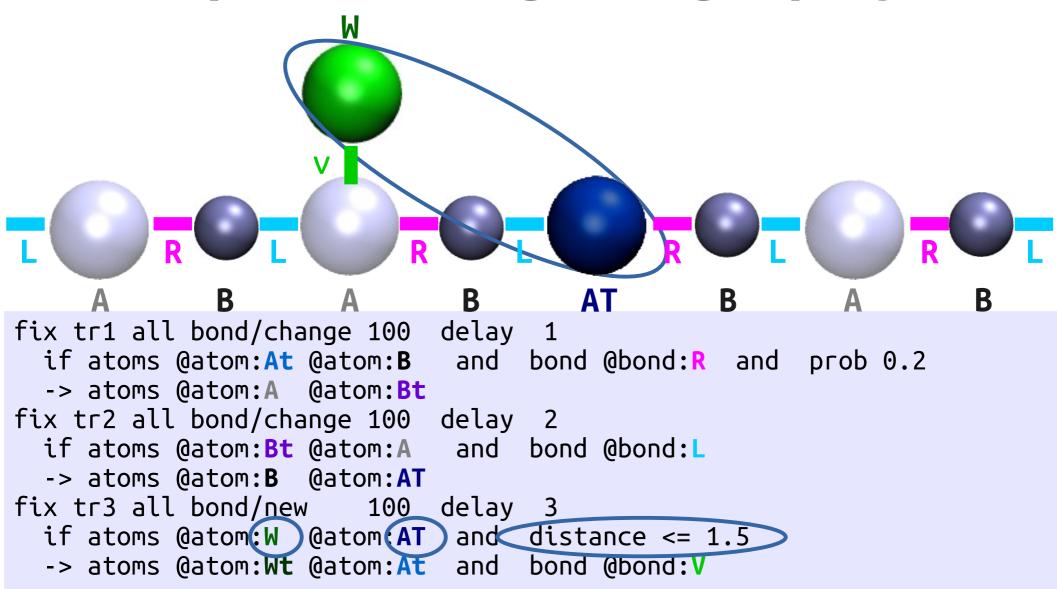


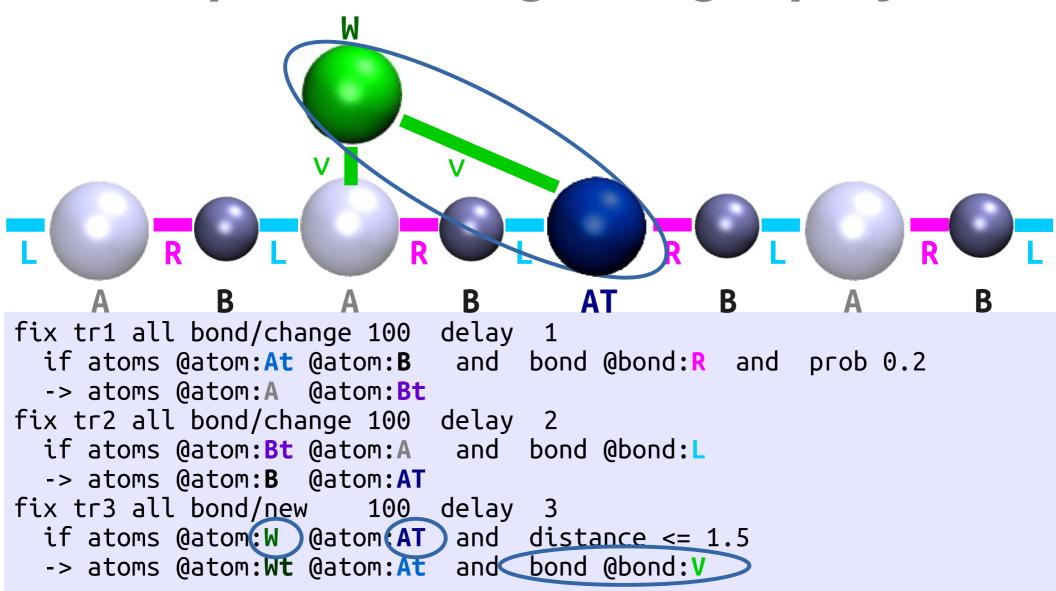


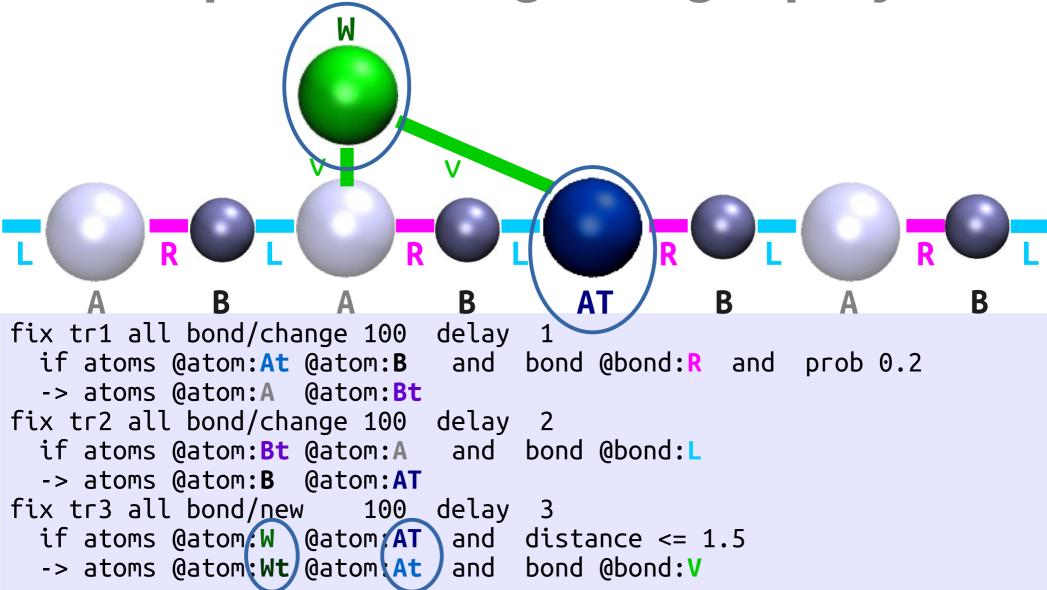


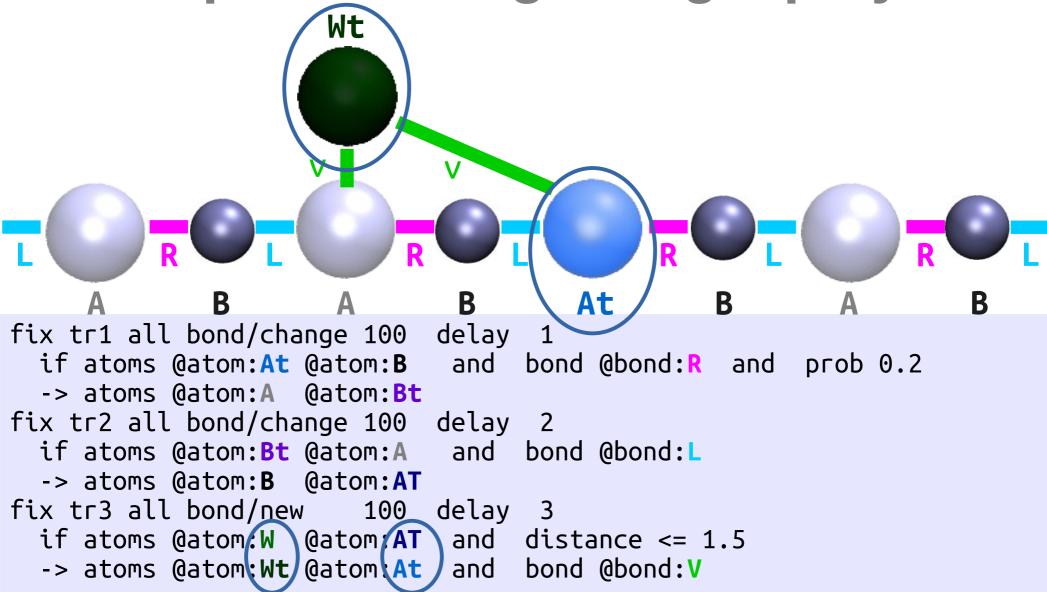


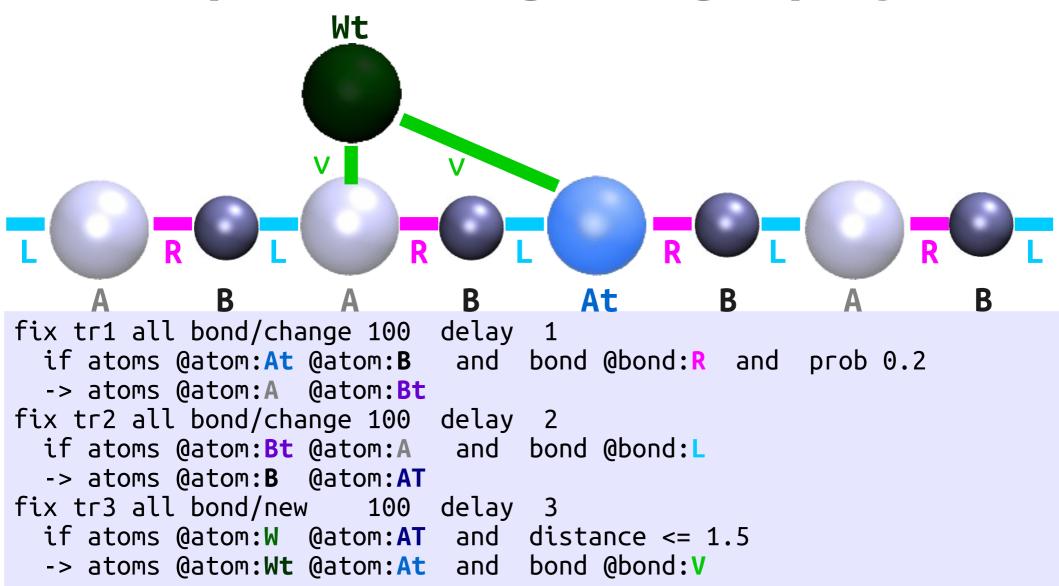


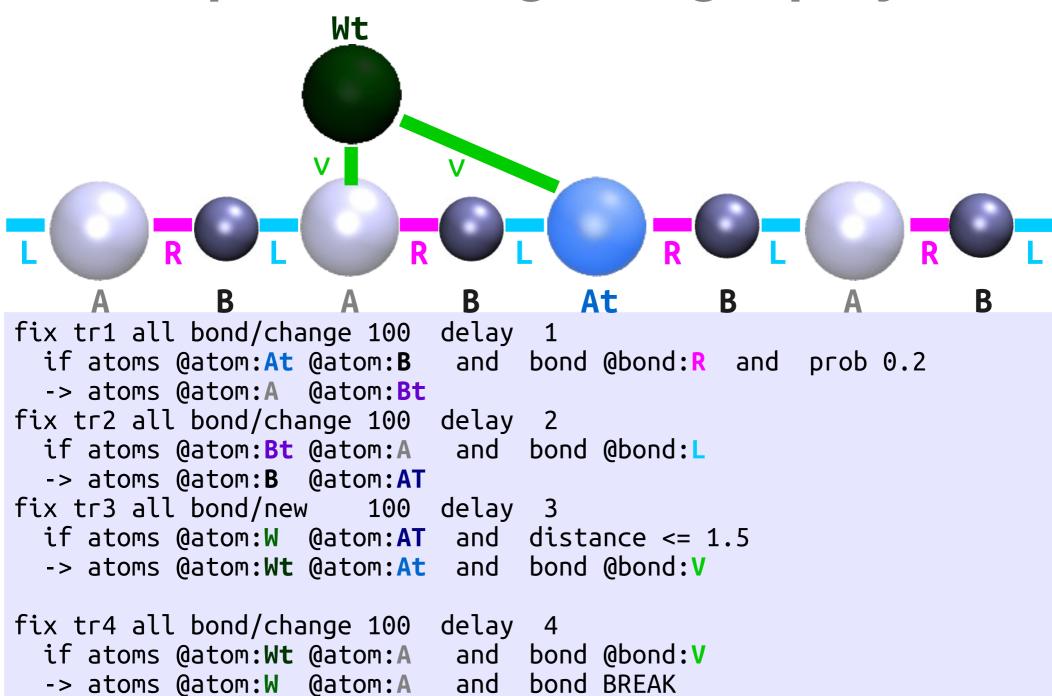


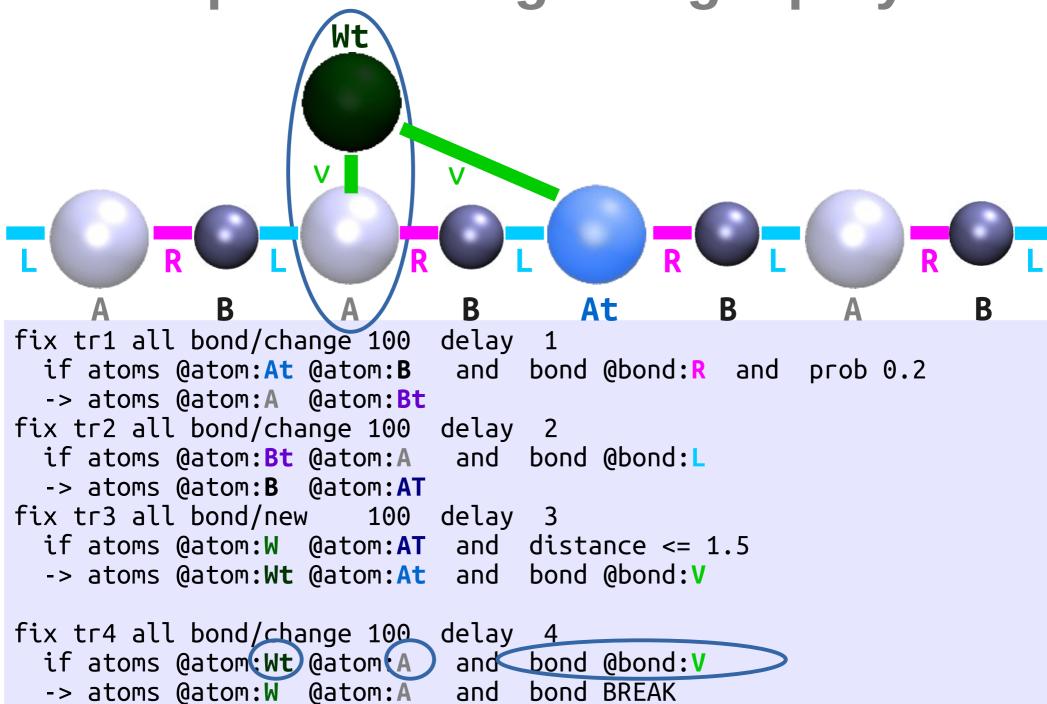


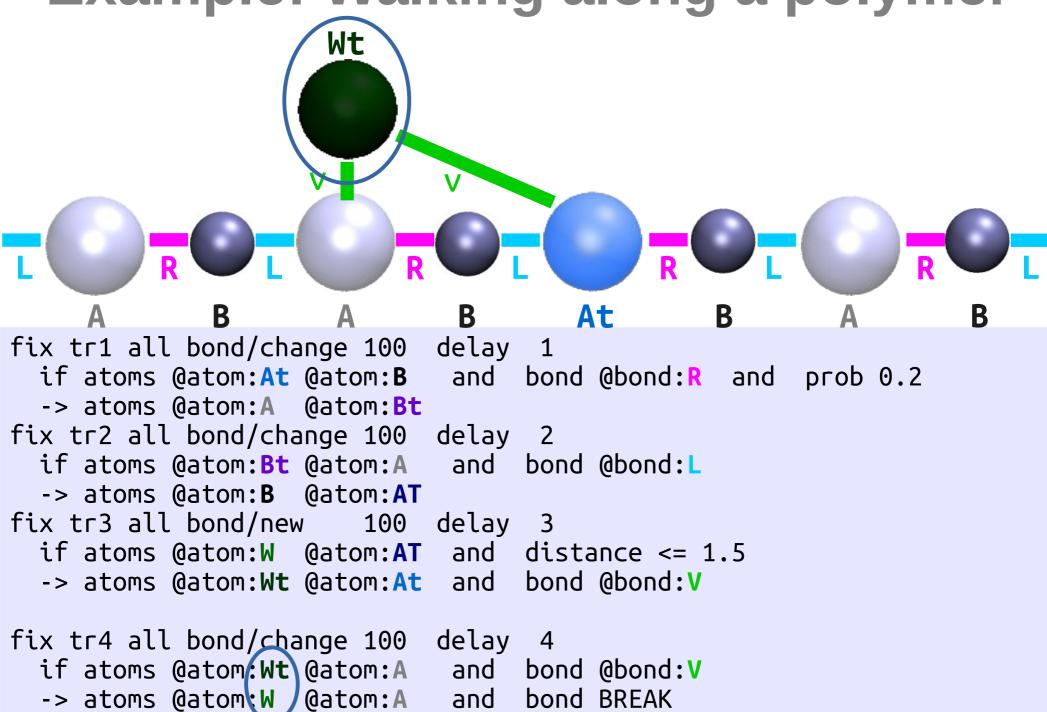


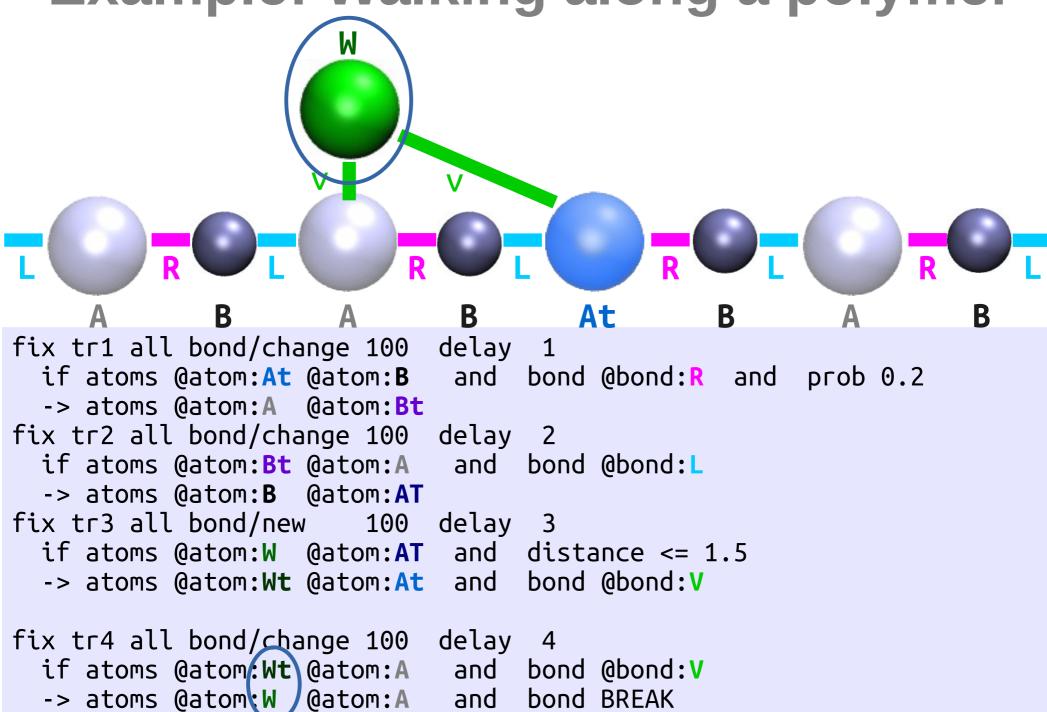


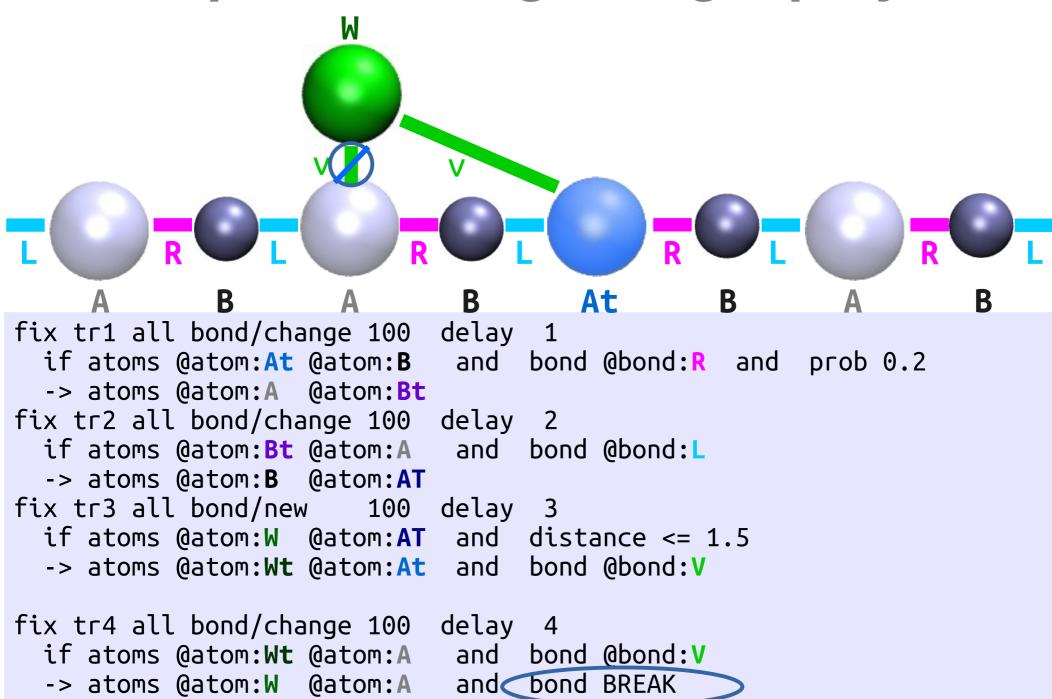


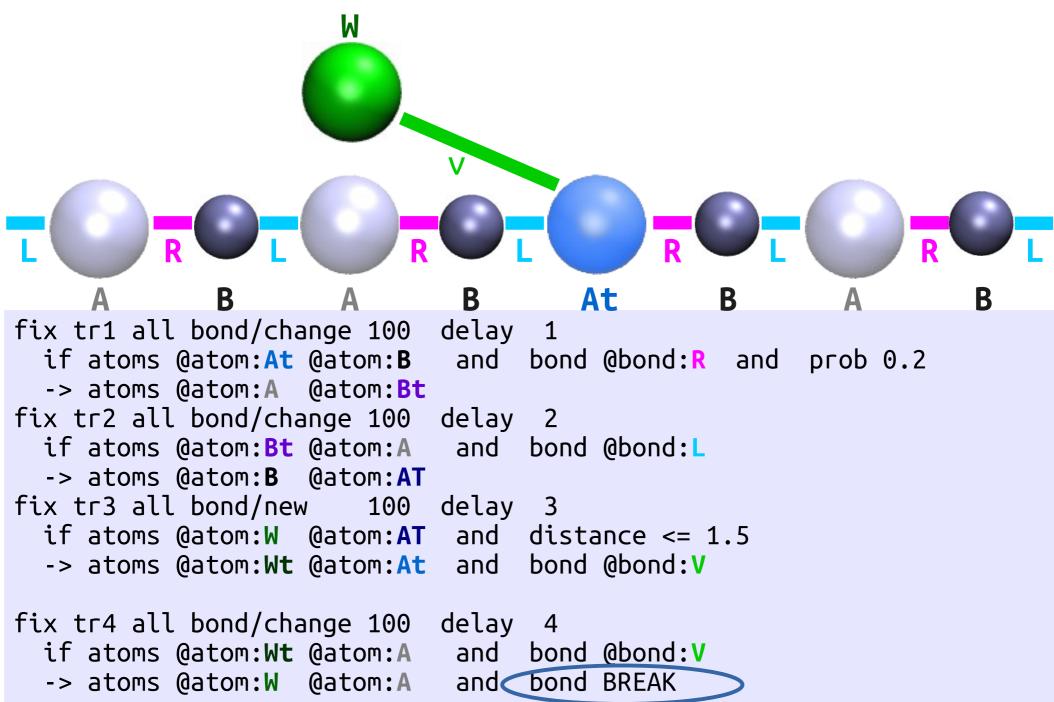


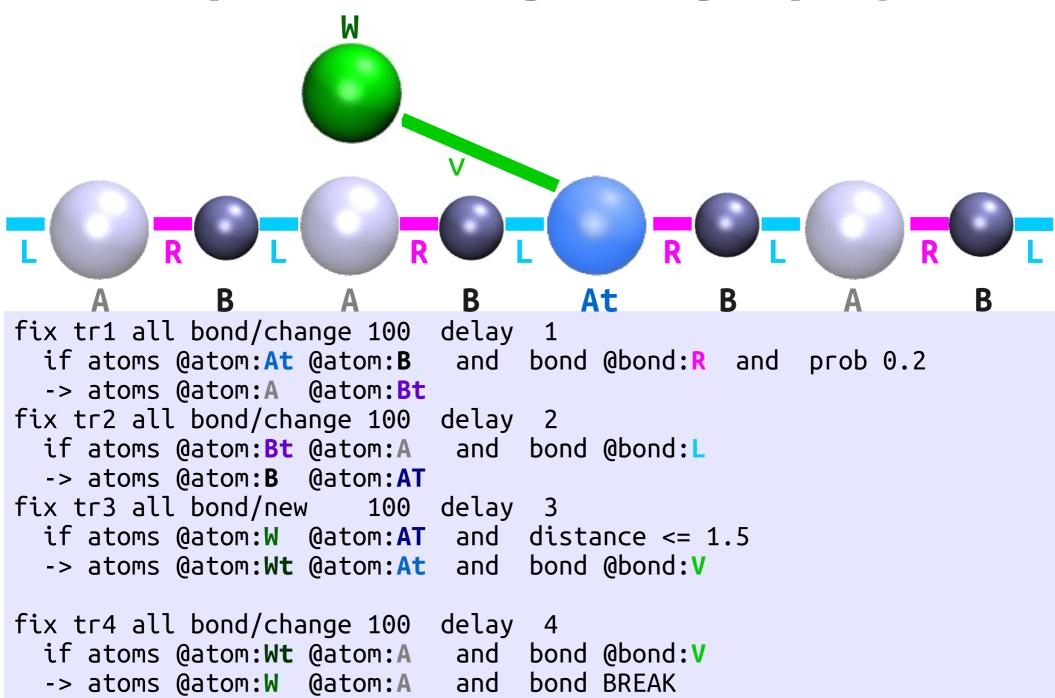


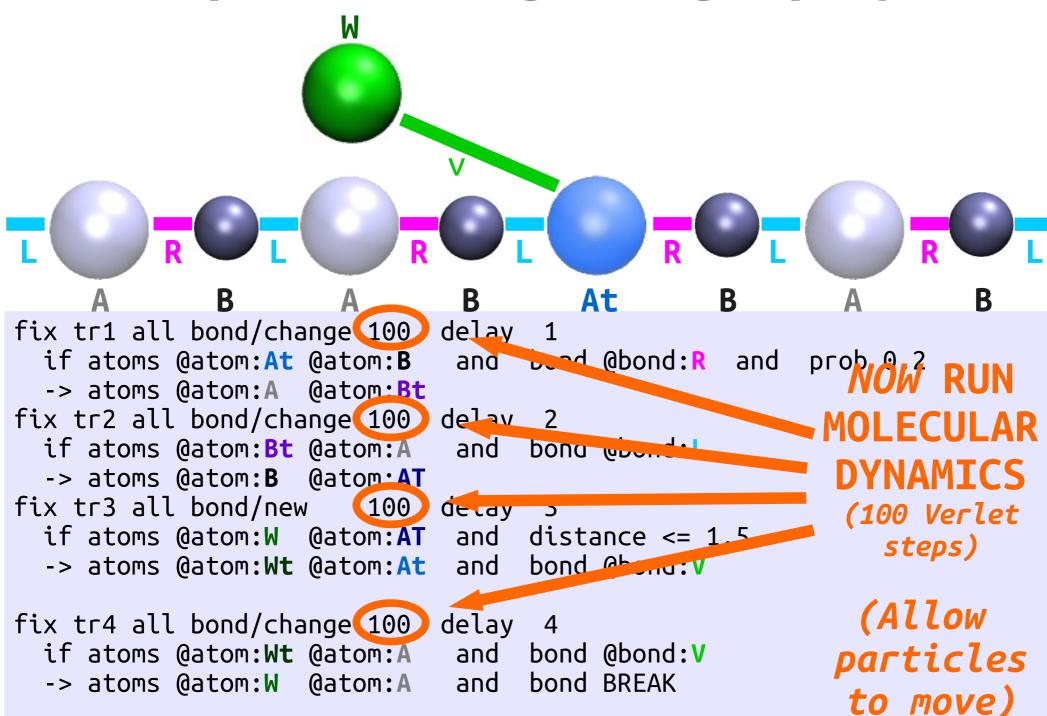


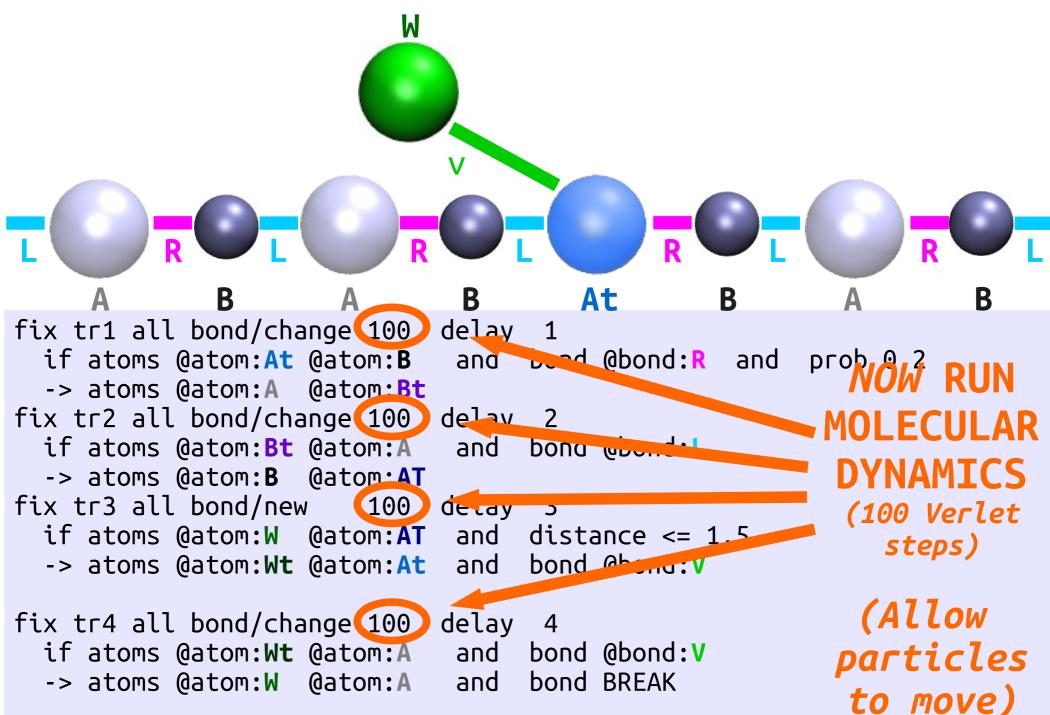


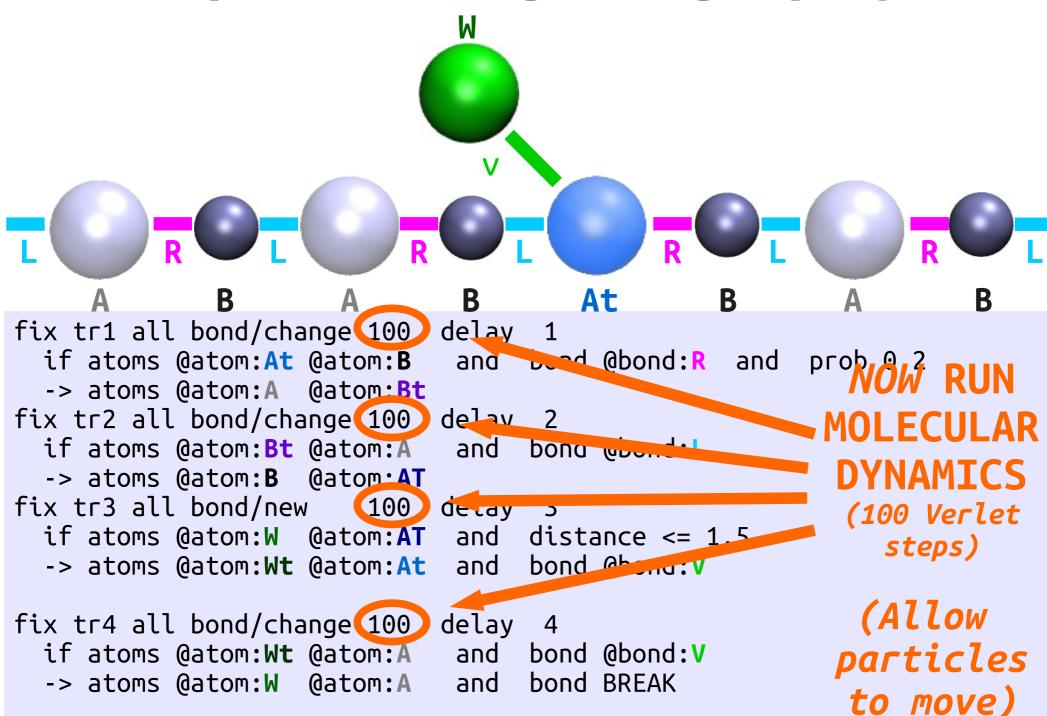


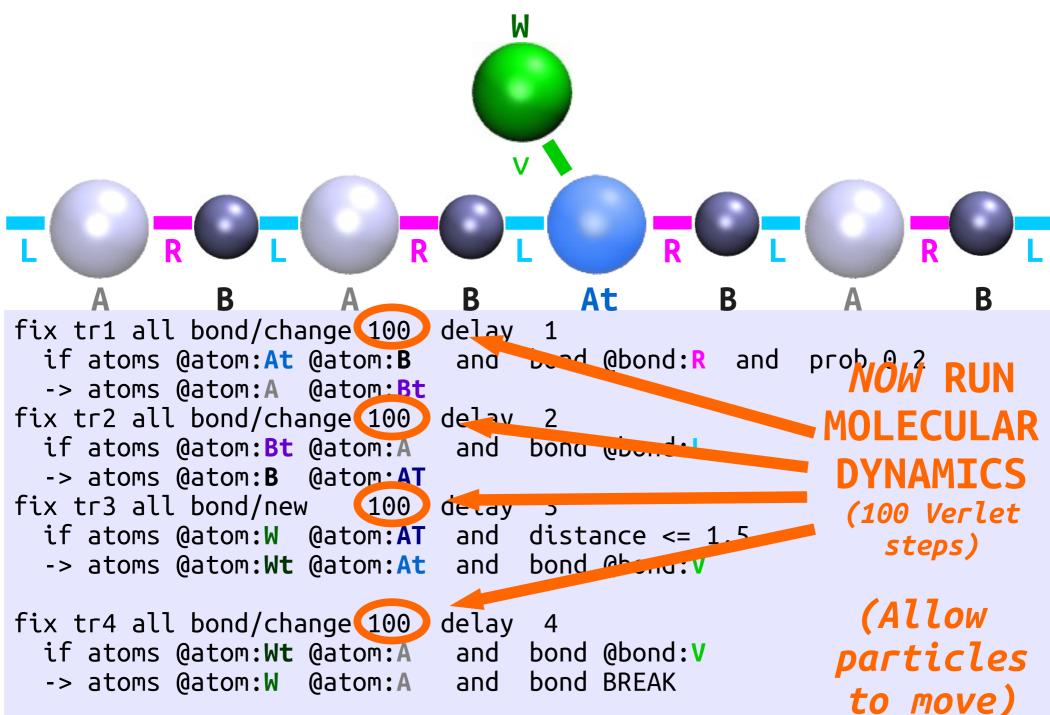


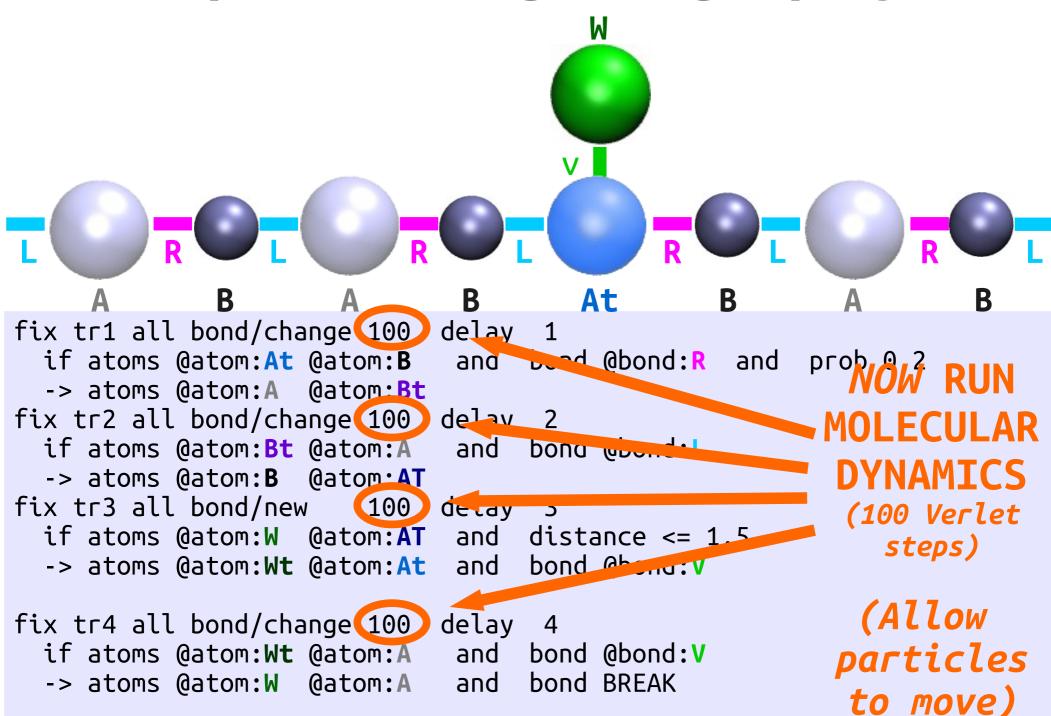


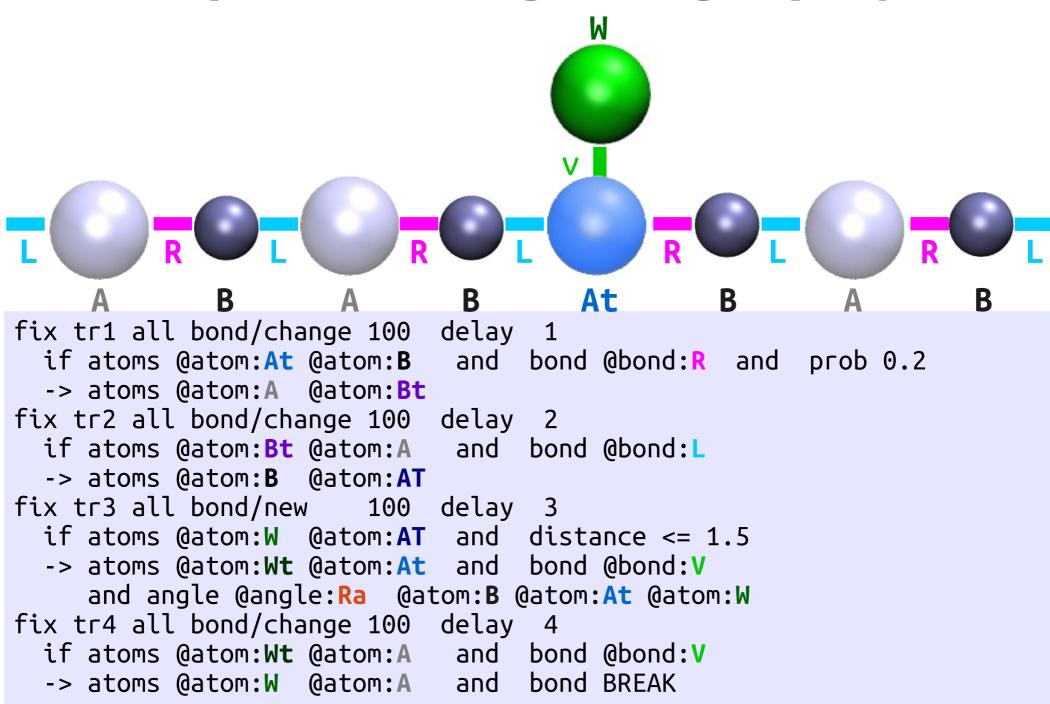










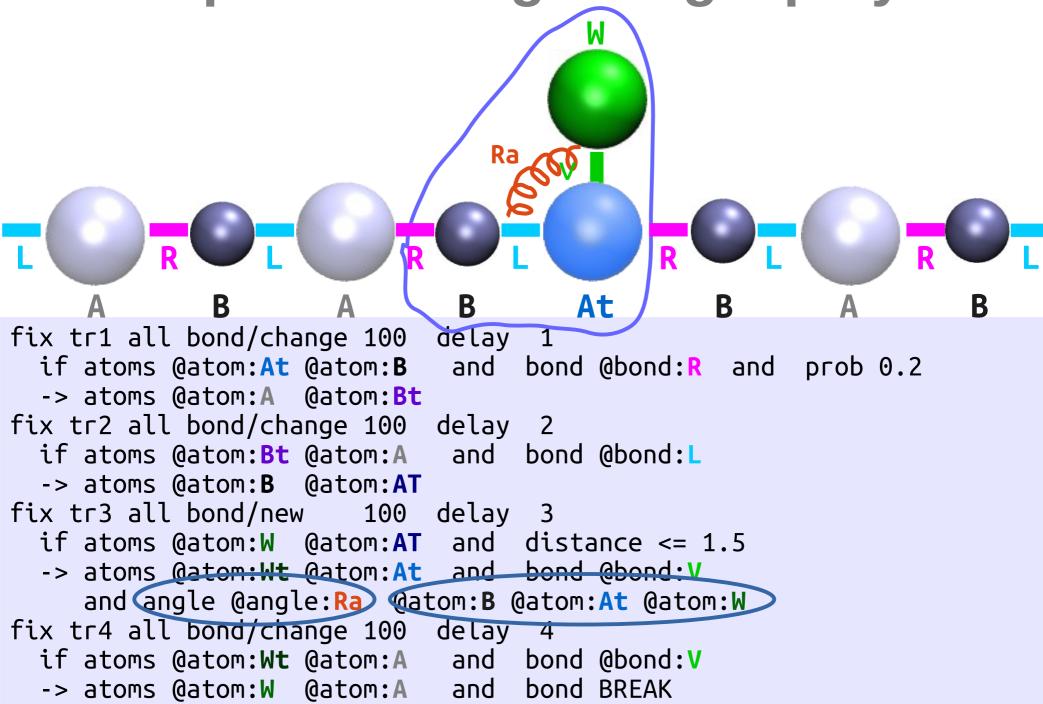


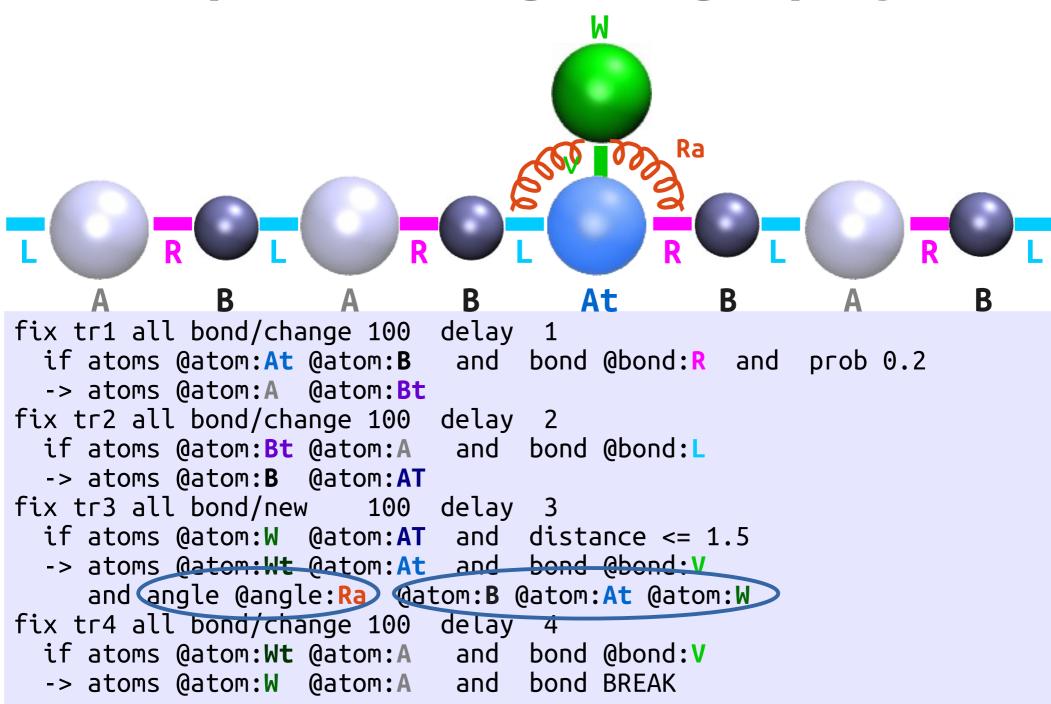
Example: Walking along a polymer B At fix tr1 all bond/change 100 delay bond @bond:R if atoms Qatom: At Qatom: B and and prob 0.2 -> atoms @atom:A @atom:Bt fix tr2 all bond/change 100 delay 2 if atoms @atom:Bt @atom:A and bond @bond:L -> atoms @atom:B @atom:AT fix tr3 all bond/new 100 delay 3 if atoms @atom:W @atom:AT and distance <= 1.5 -> atoms @atom:Wt @atom:At and bond @bond:V and angle @angle:Ra @atom:B @atom:At @atom:W fix tr4 all bond/change 100 delay 4 if atoms @atom:Wt @atom:A and bond @bond:V

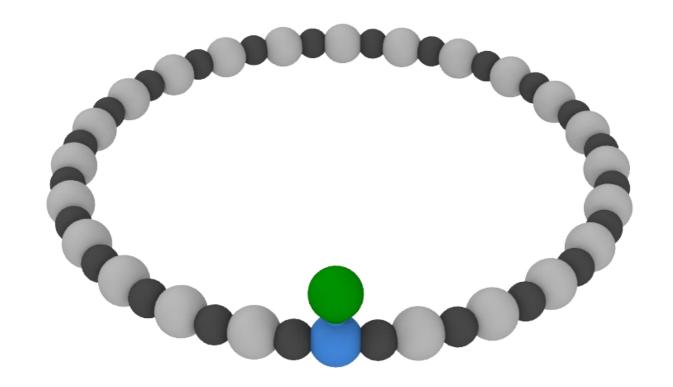
and

bond BREAK

-> atoms @atom:W @atom:A

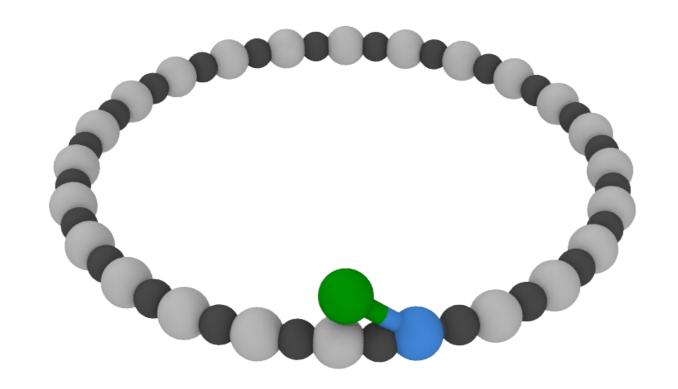




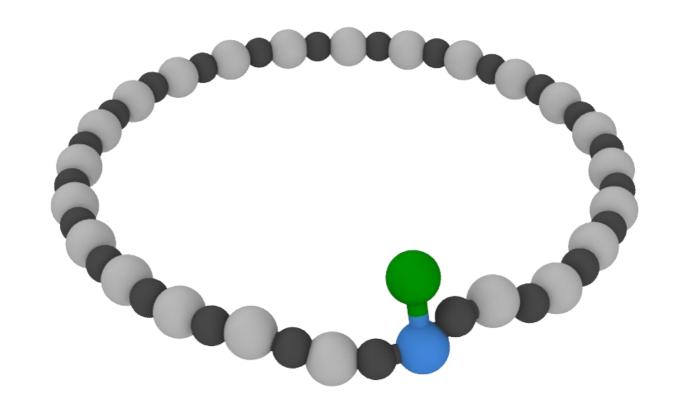


#### SEE:

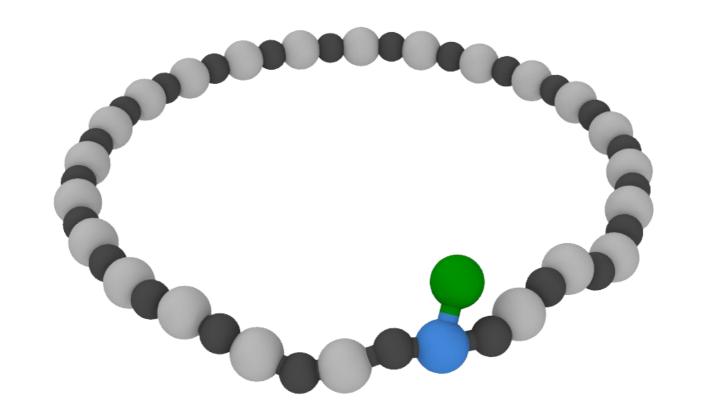
https://www.youtube.com/watch?v=QO4LbHGAgxU



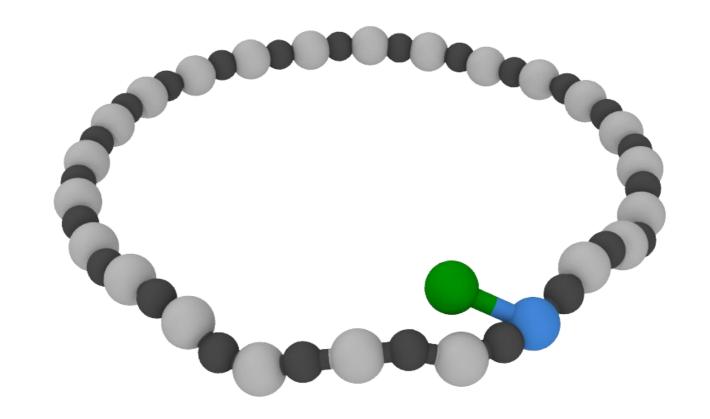
#### SEE:



#### SEE:



#### SEE:



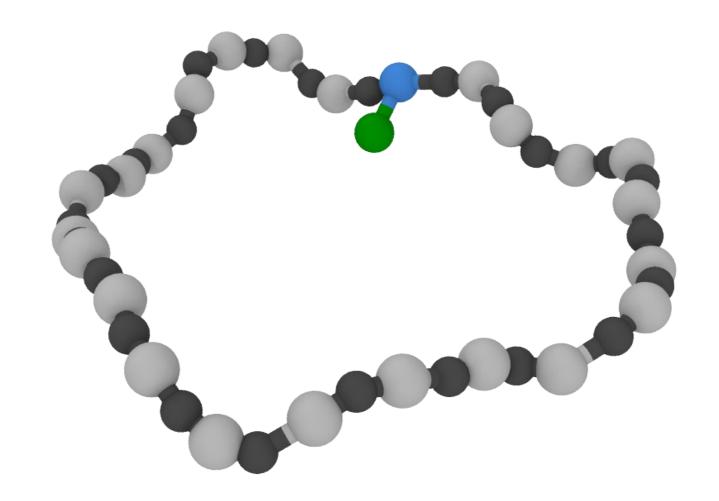
#### SEE:



SEE:

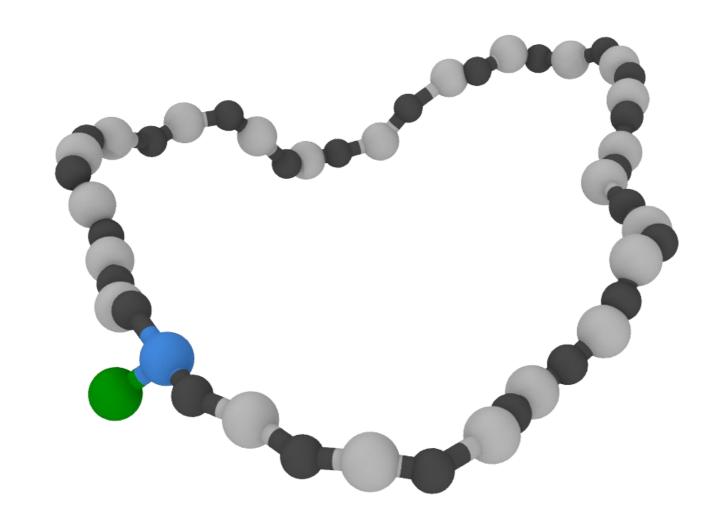


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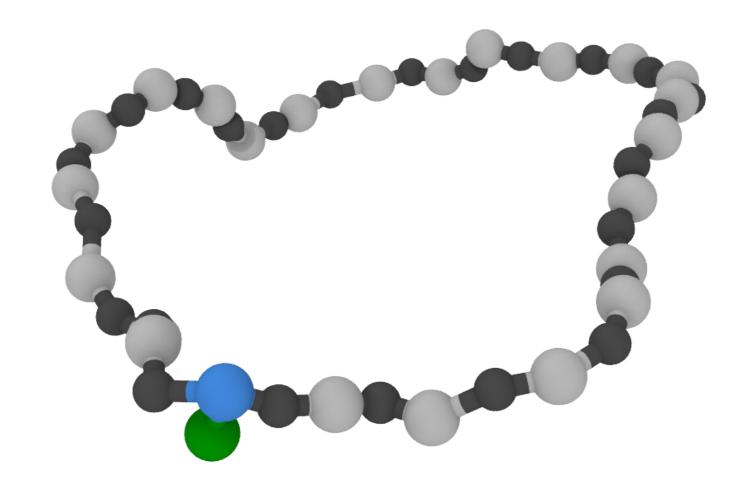


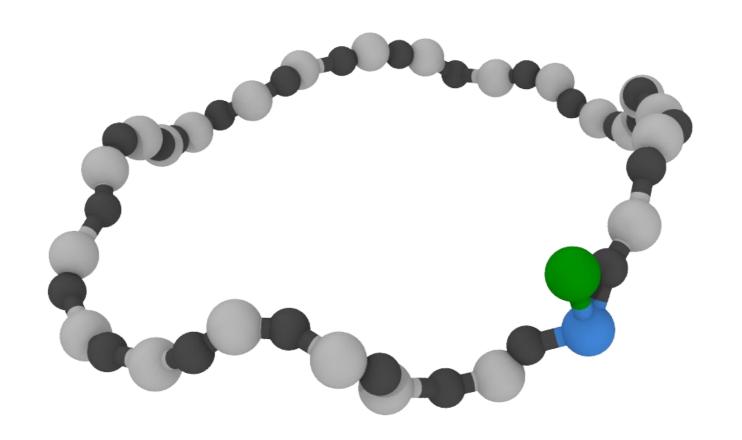
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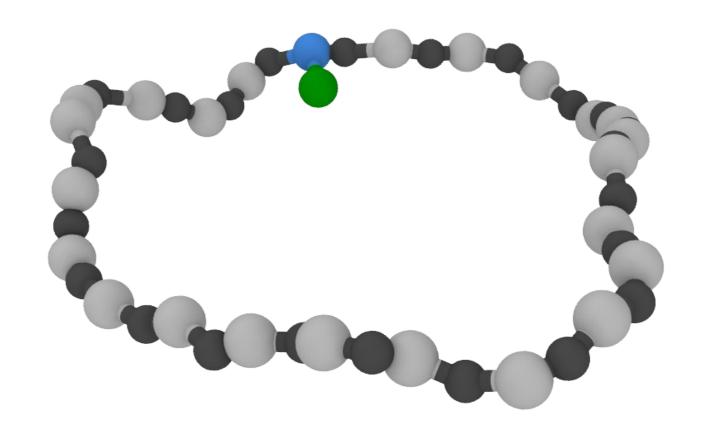


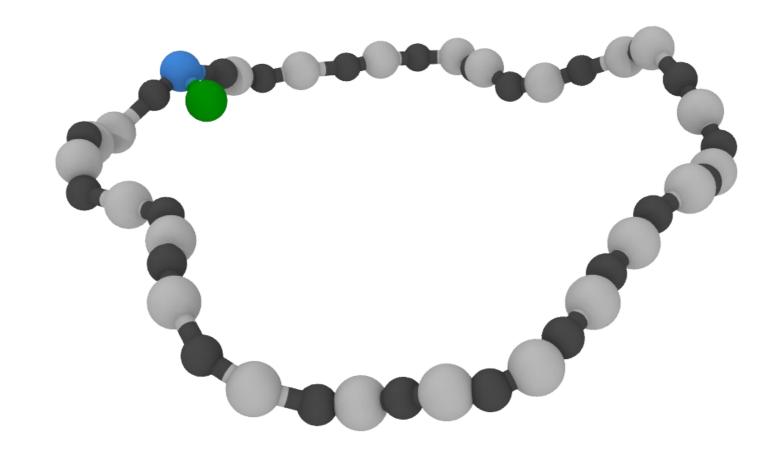


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