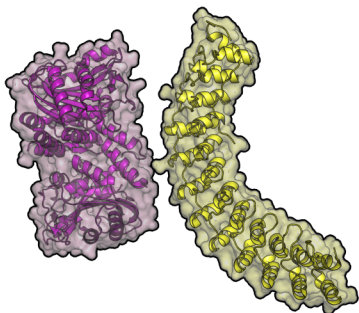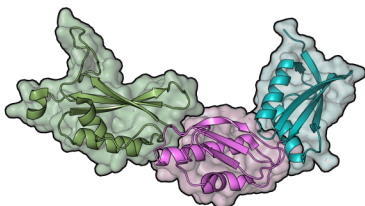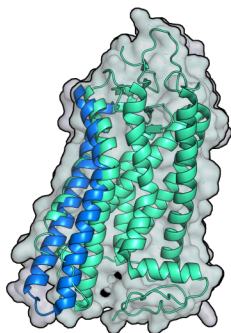Two-domain rigid body docking

Multi-domain rigid body docking

Conformational change modeling

De novo fold prediction

```
>T4 Lysozyme
MNIFEMLRIDEGLRLKI...
```

Modeling complexity