

A large, semi-transparent graphic on the left side of the slide. It consists of a large orange circle with a white ring inside, similar to the SPARC logo. This graphic is overlaid on a background of numerous small, colorful dots in shades of orange, red, purple, and blue, which appear to be a microscopic view of cells or a simulation of a biological process.

Informatics and Computer Simulations: Experience of SPARC

23rd May 2019, Budapest

Computational Simulations of Pharmaceutical Formulations: A Case study



1. Components
2. Parameters for analysis
3. Evolution from packing to nanoparticle formation
4. Snapshots of nanoparticle formation
5. Analysis
6. Summary

System input

Total number of atoms 327376

Ensemble: NPT

Total simulation time: 200ns

Formulation components (considering one half of original system):

Drug molecule	30
Fatty acid	15
PEG	456
Cholesterol	4
PVP	6
Ethanol	543
Water	99270
Sodium ions for neutral pH	4
TOTAL	100,327

Parameters considered for analysis are:

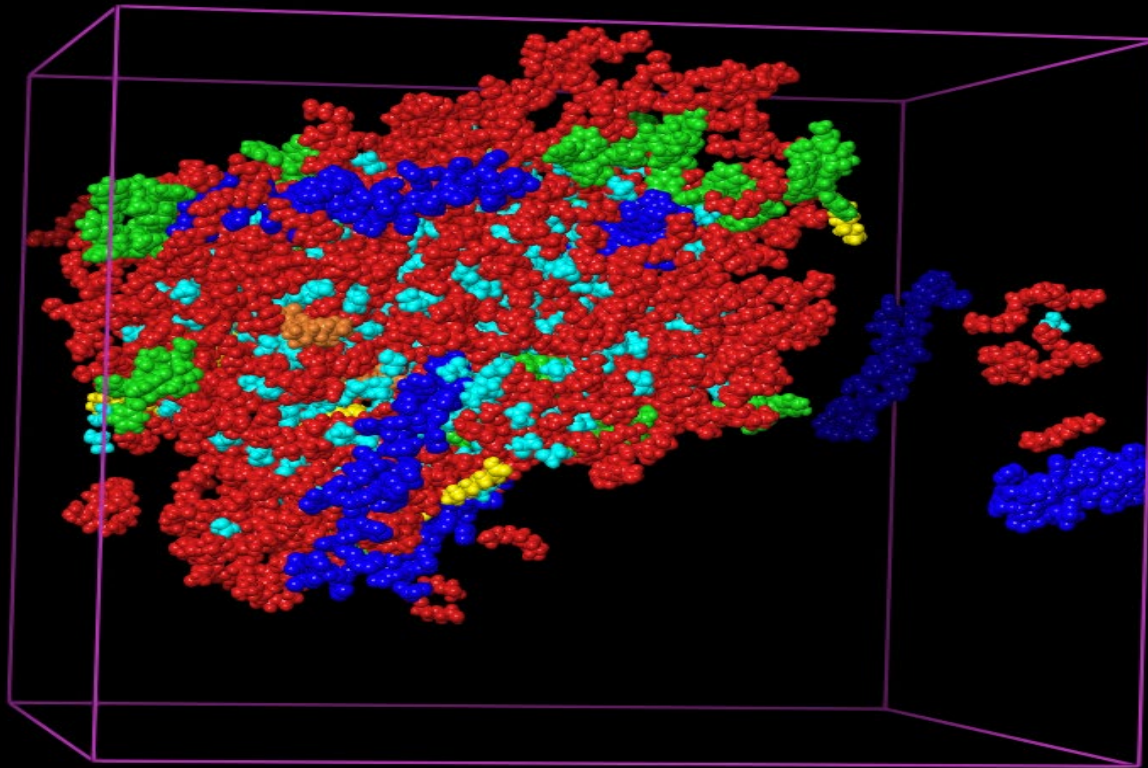
H-bond analysis

RMSD

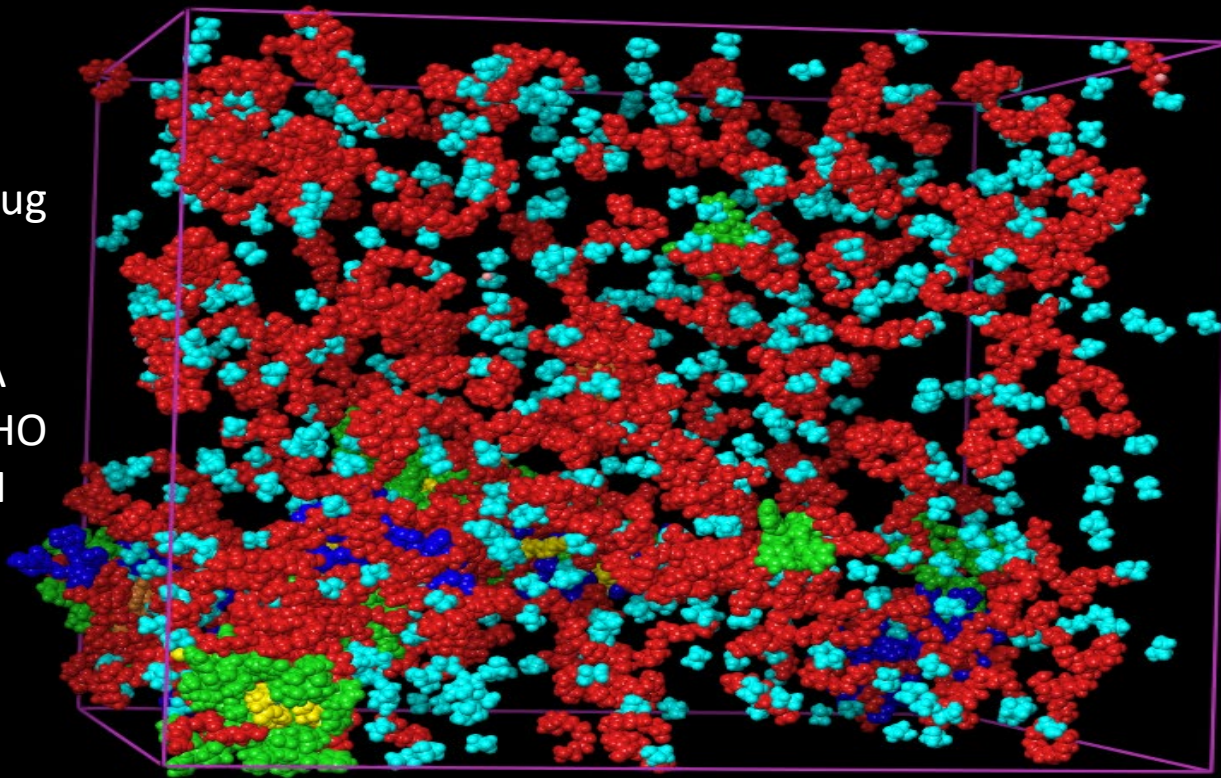
Potential energy

Simulating Drug Molecule...

Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO
Cyan=EtOH

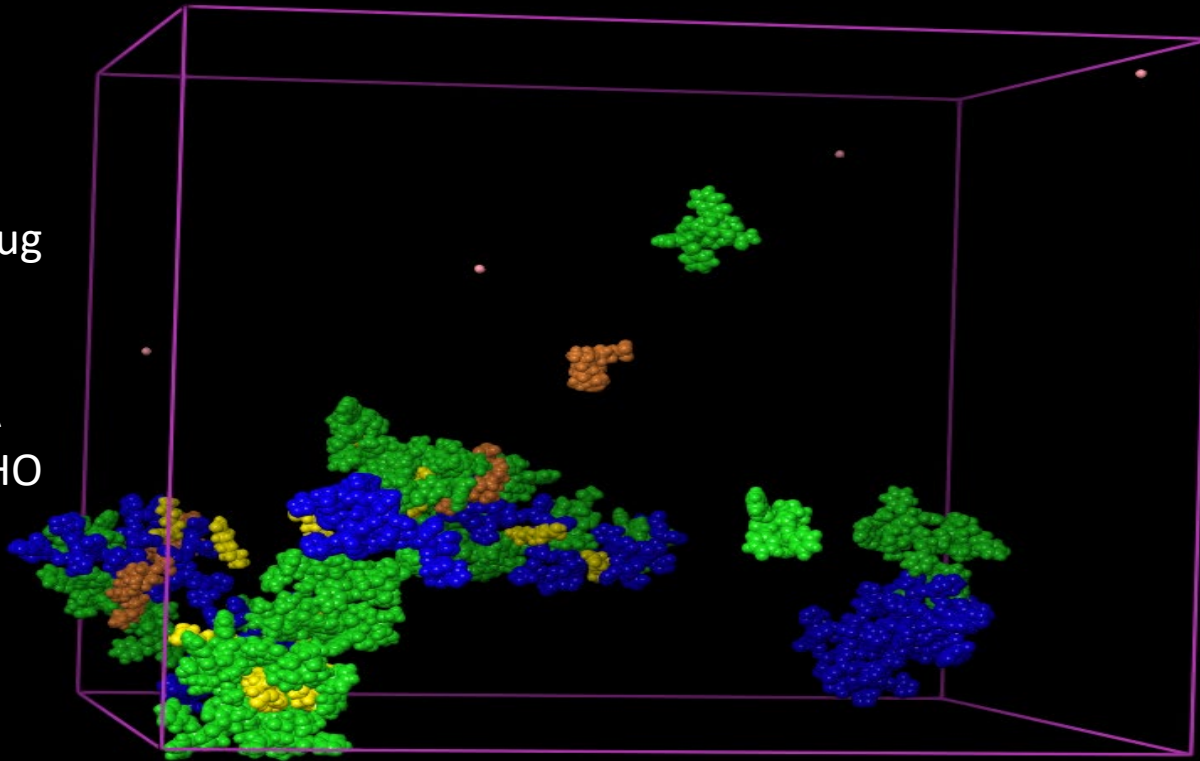


Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO
Cyan=EtOH



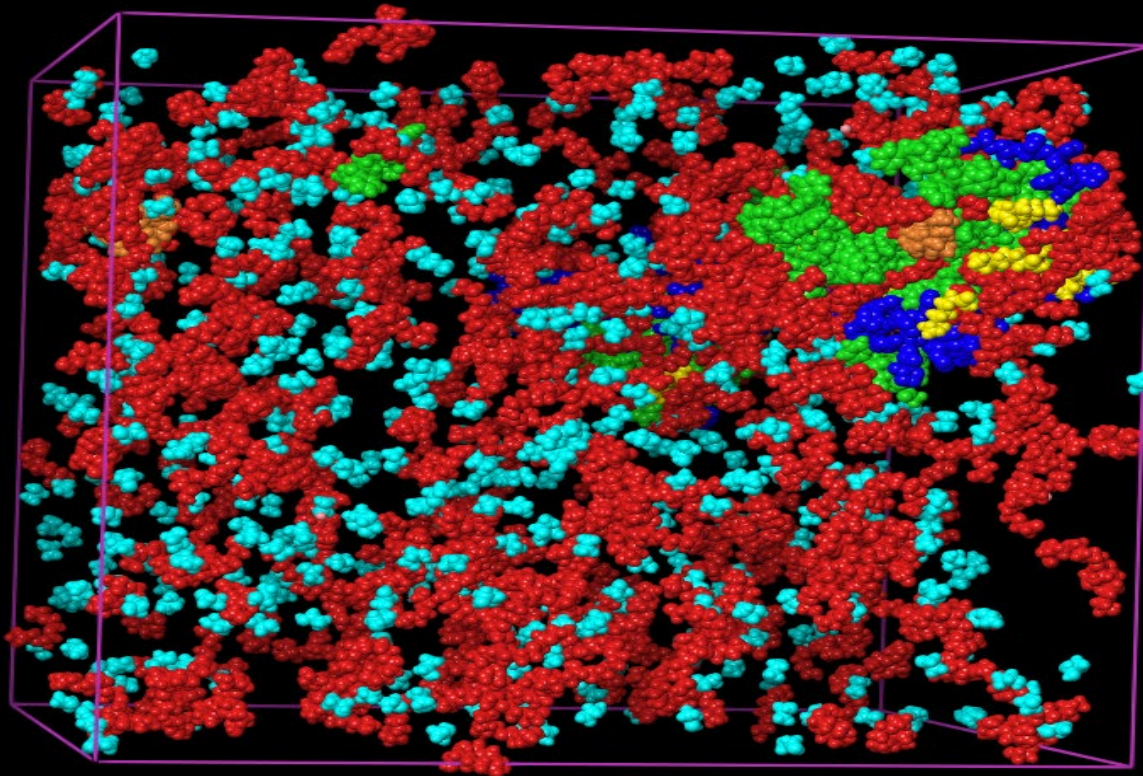
T=20 ns

Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO



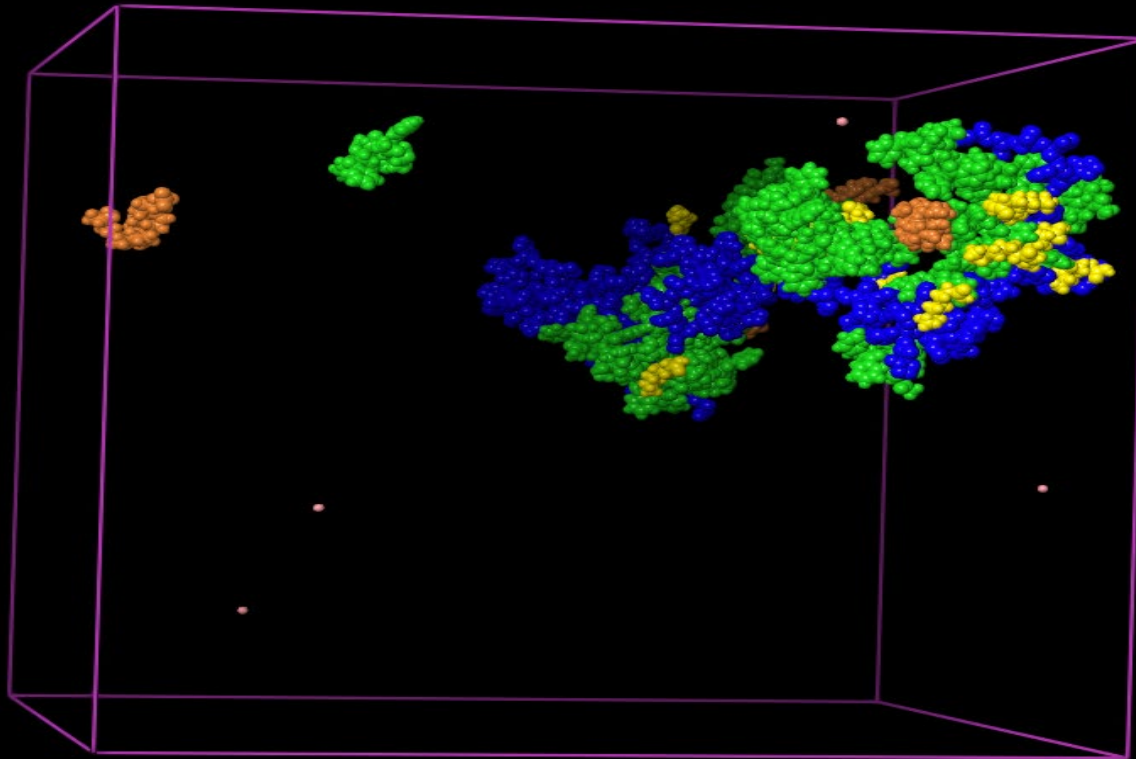
T=40 ns

Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO
Cyan=EtOH



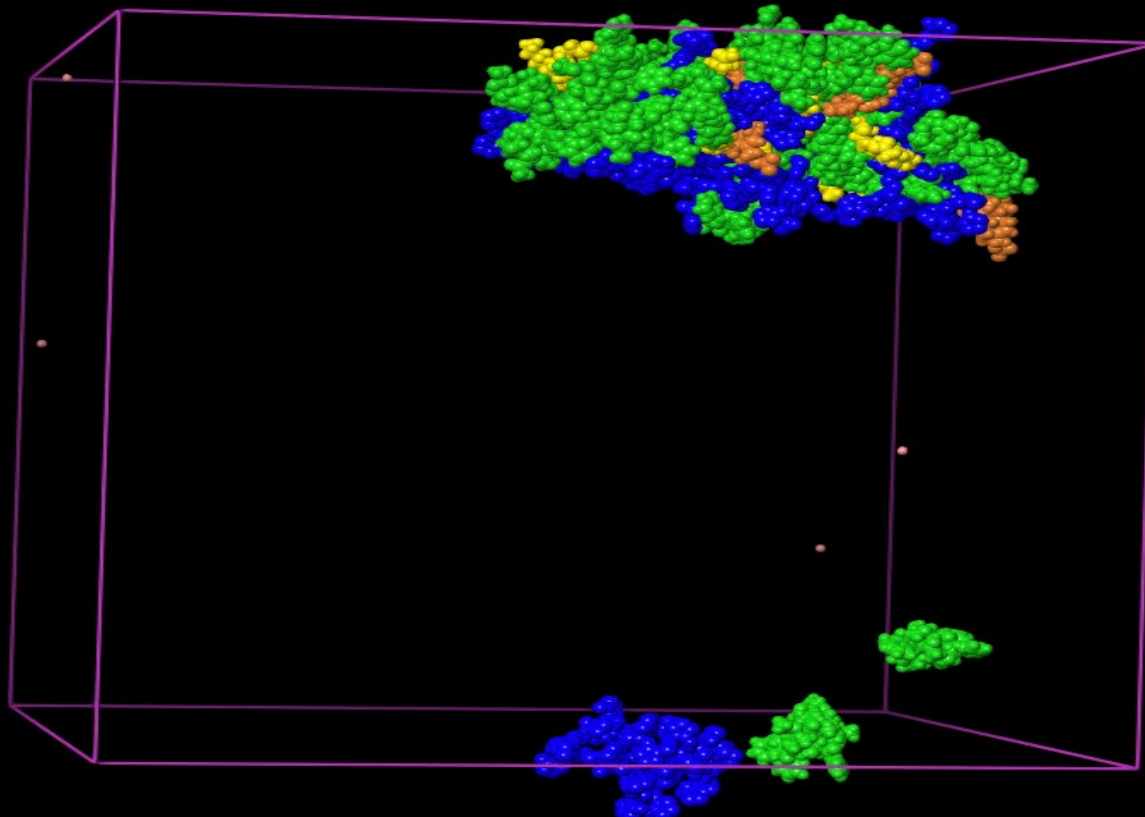
T=40 ns

Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO
Cyan=EtOH



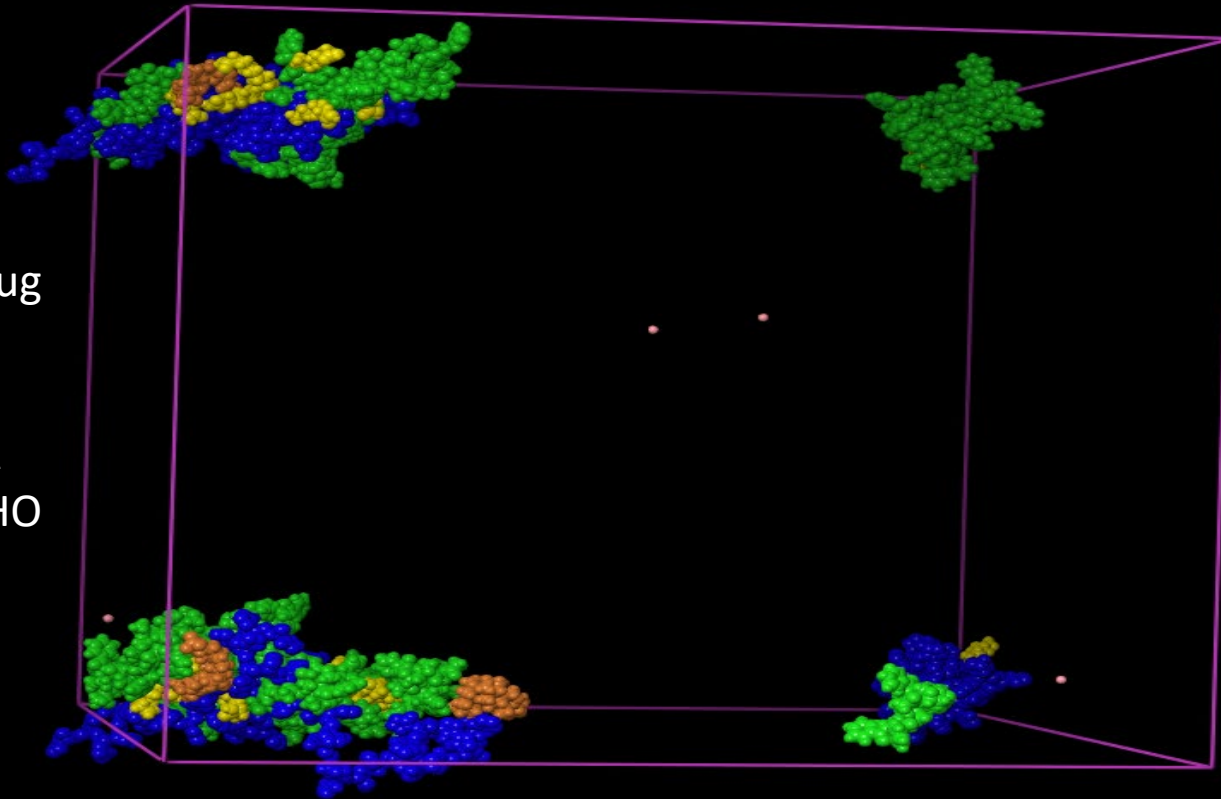
T=60 ns

Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO
Cyan=EtOH



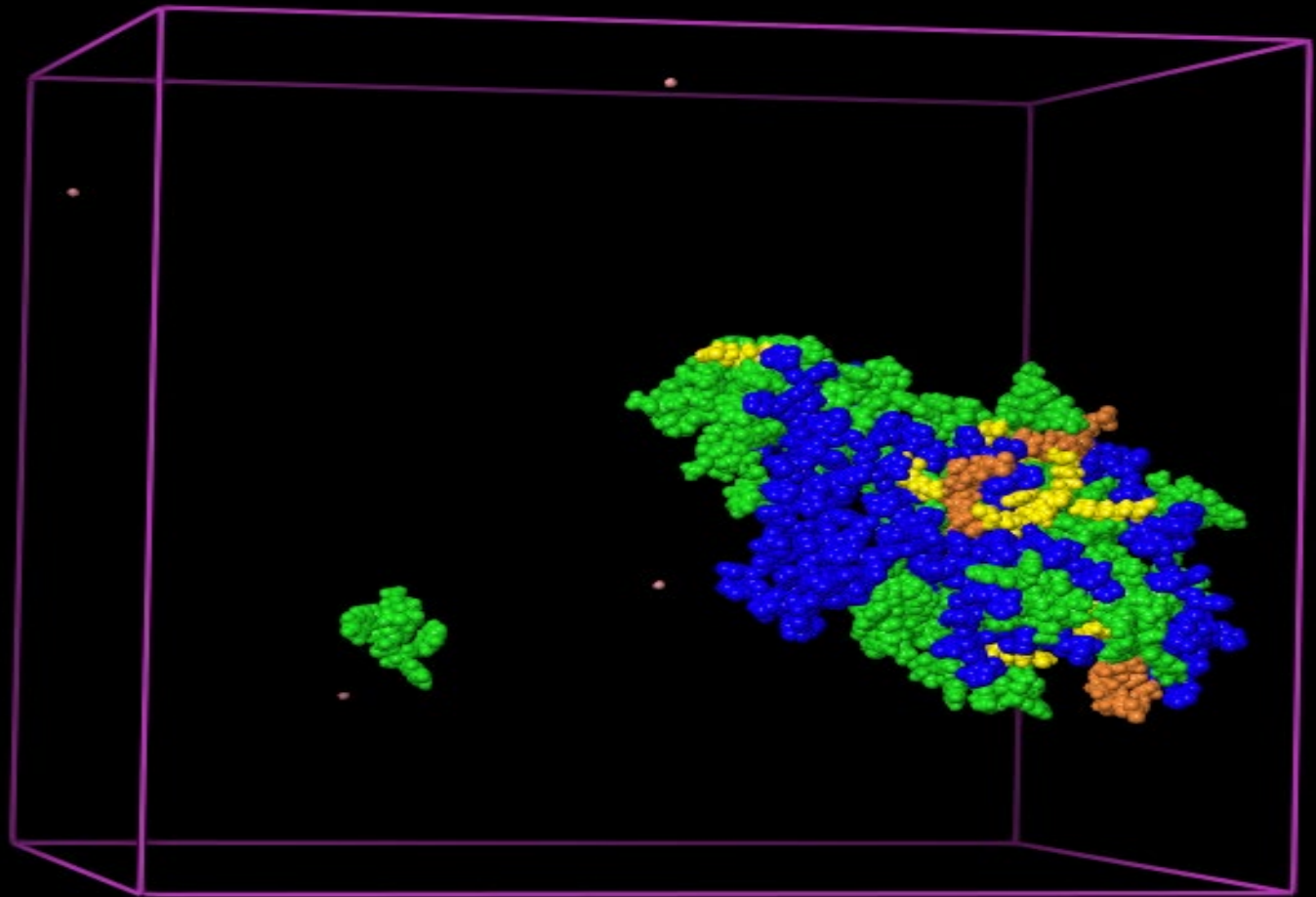
T=80 ns

Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO
Cyan=EtOH



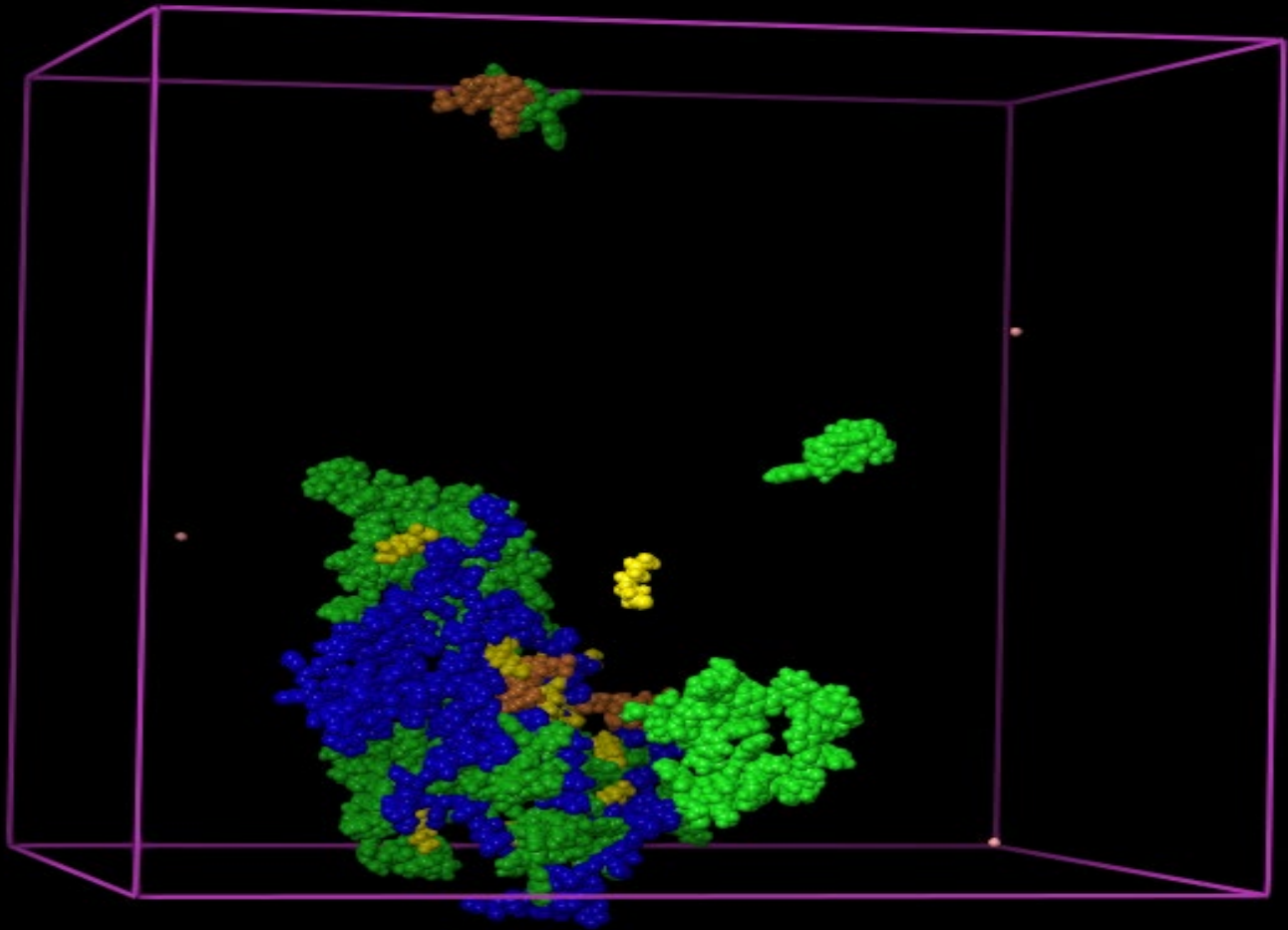
T=100 ns

Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO
Cyan=EtOH



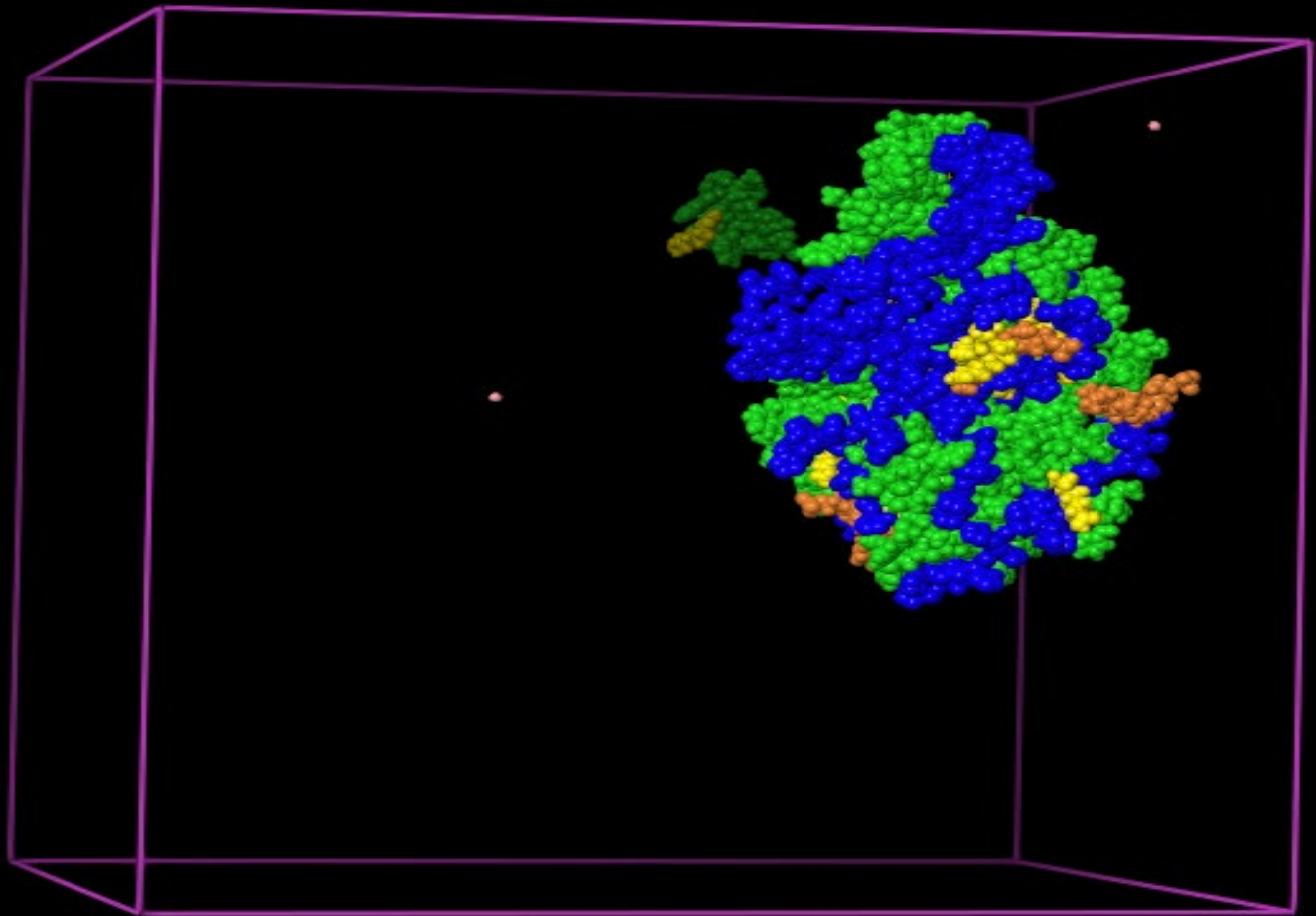
T=120 ns

Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO
Cyan=EtOH



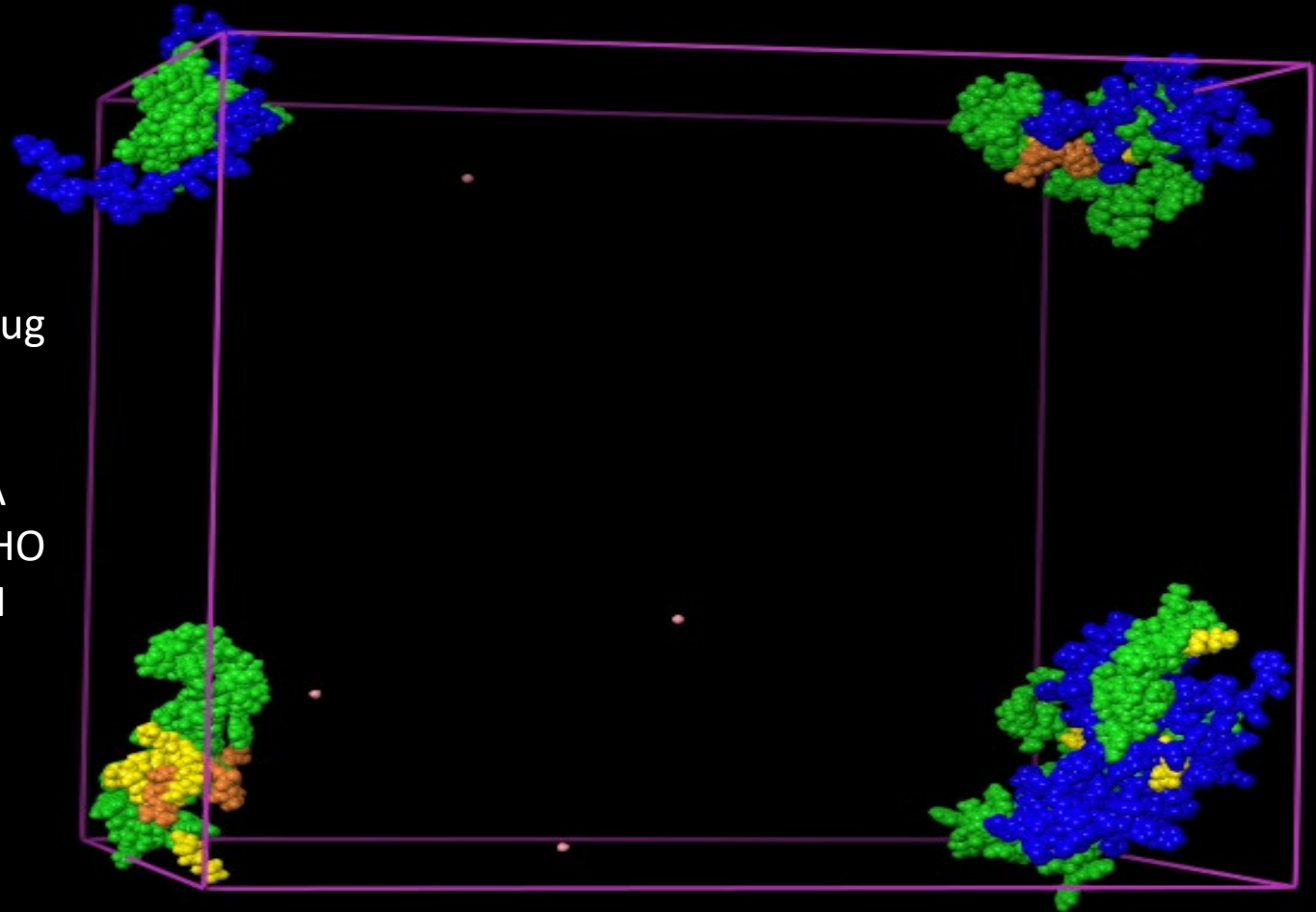
T=140 ns

Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO
Cyan=EtOH



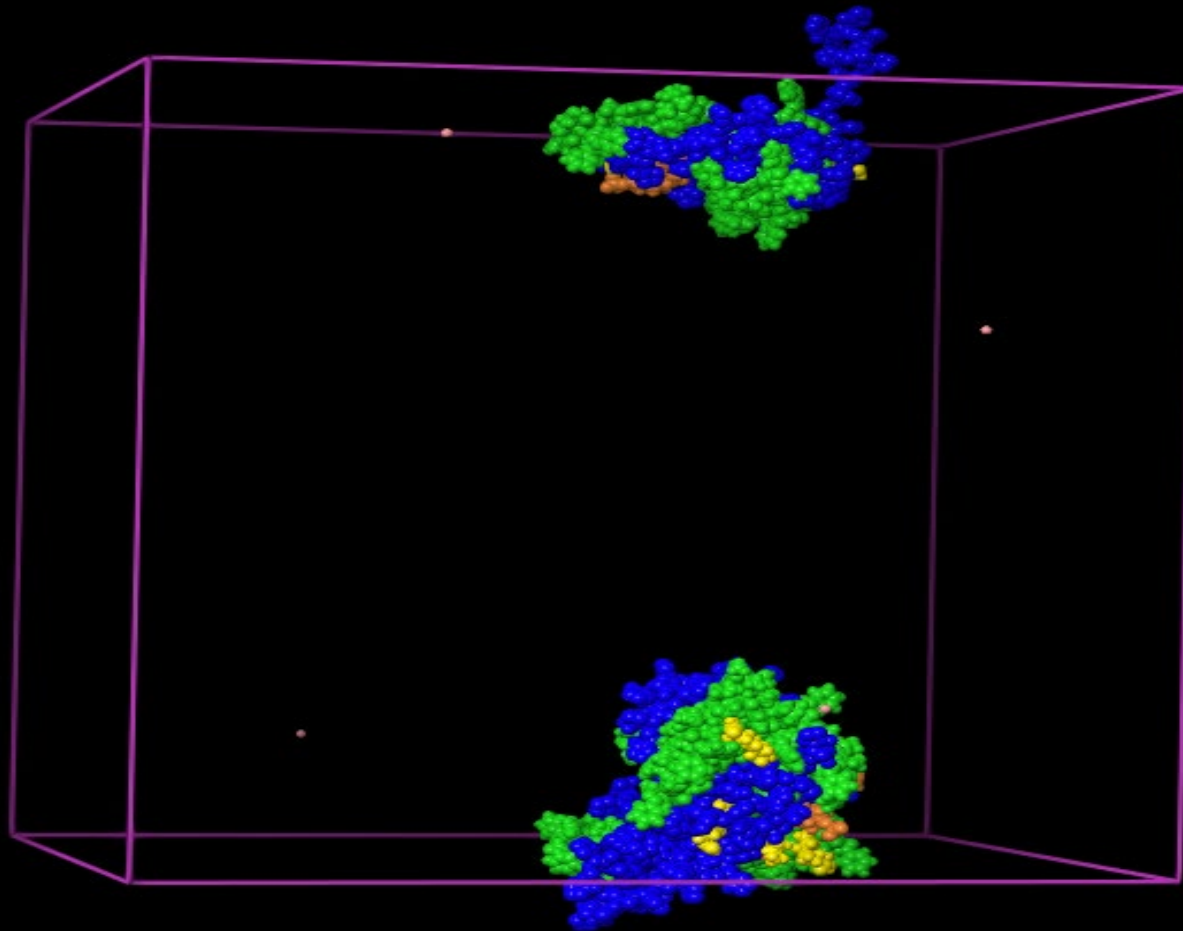
T=160 ns

Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO
Cyan=EtOH



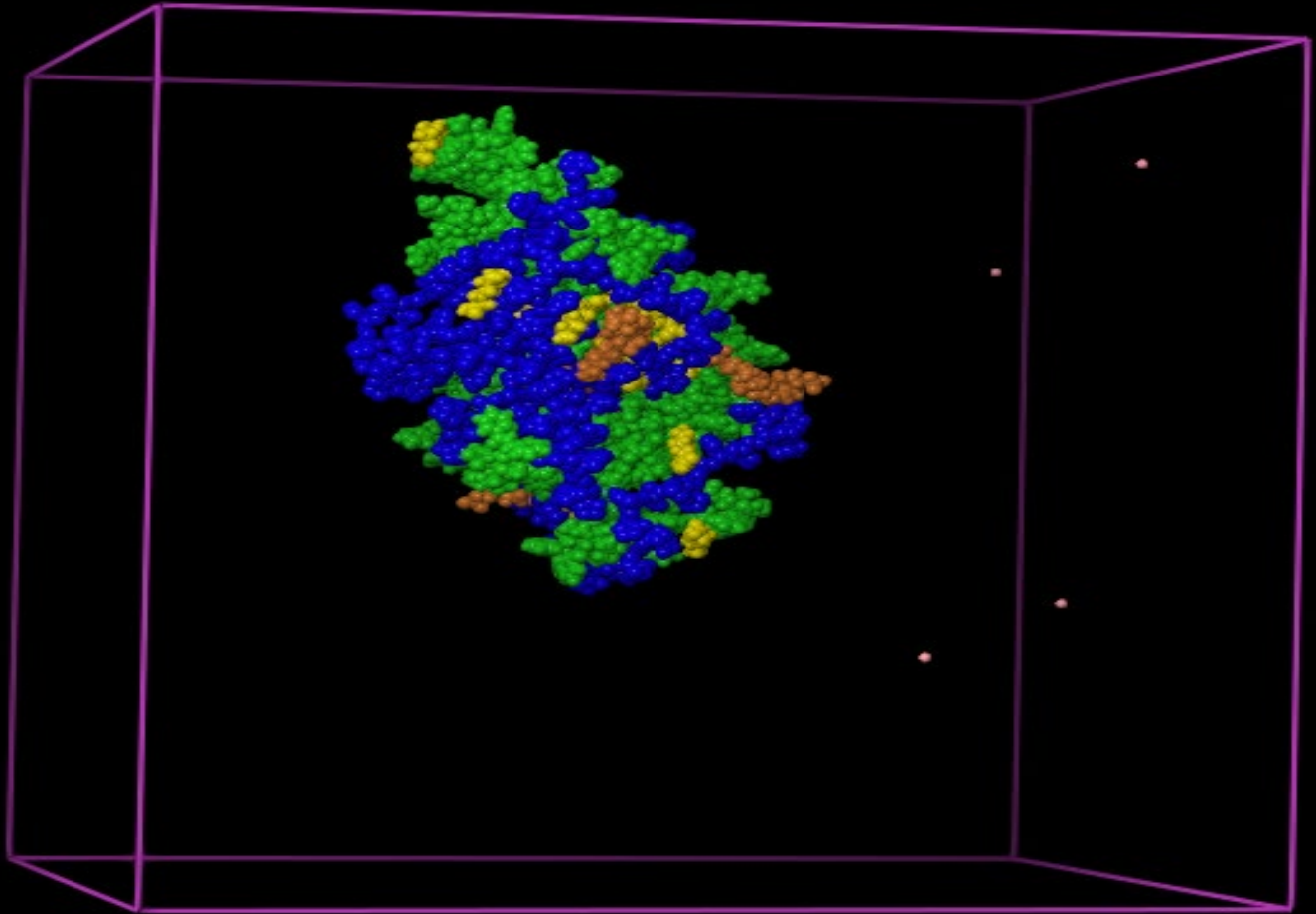
T=180 ns

Green = Drug
Blue = PVP
Red = PEG
Yellow = FA
Brown = CHO
Cyan=EtOH

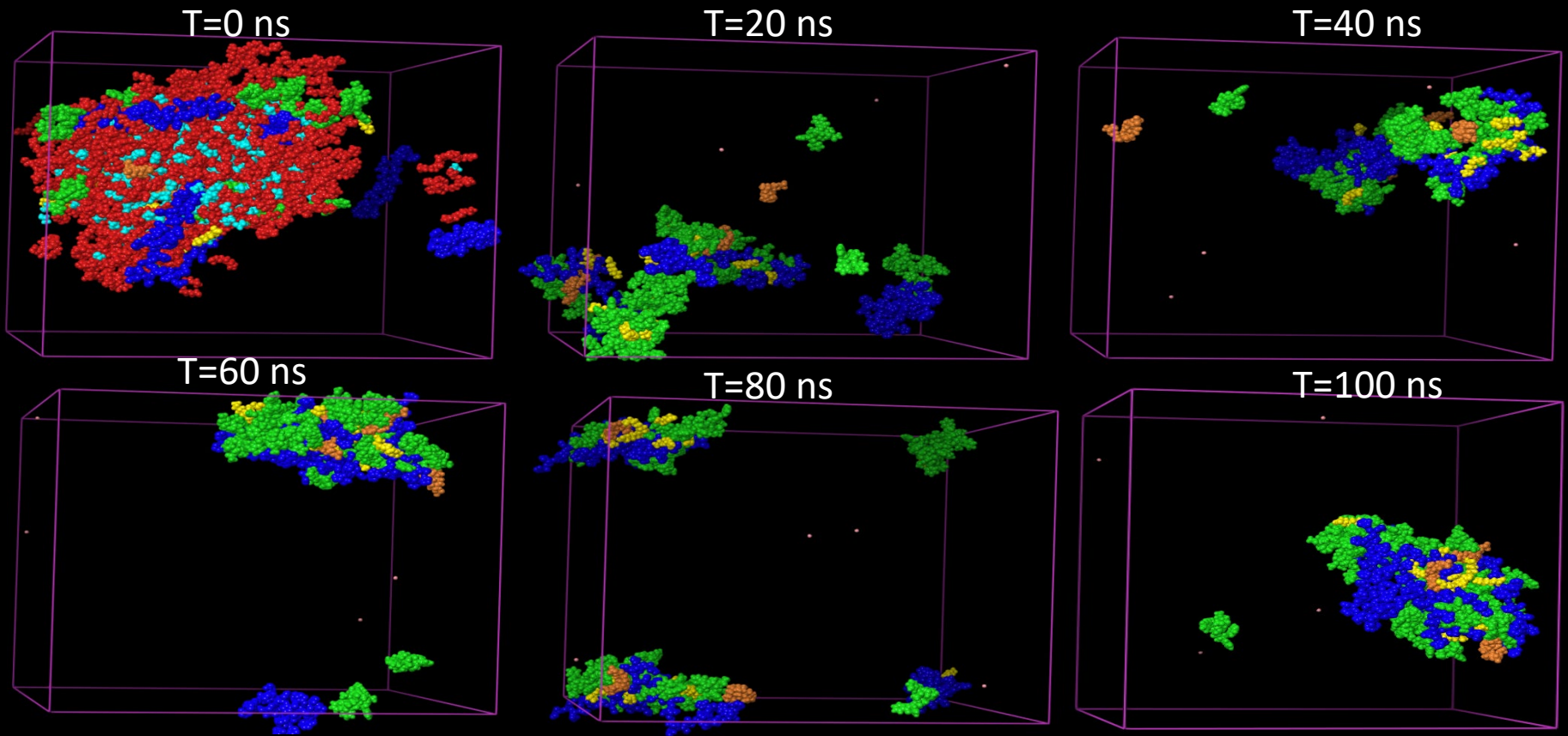


T=200 ns

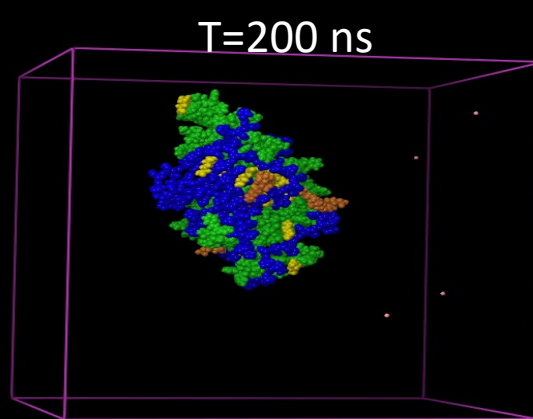
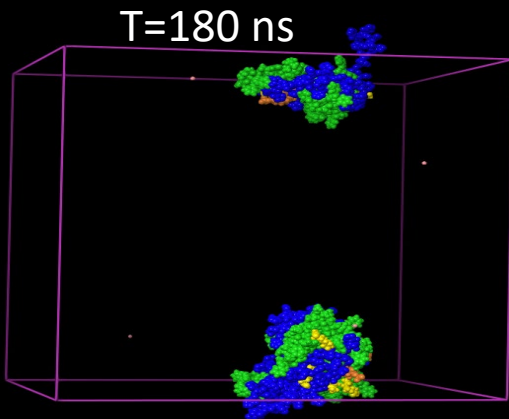
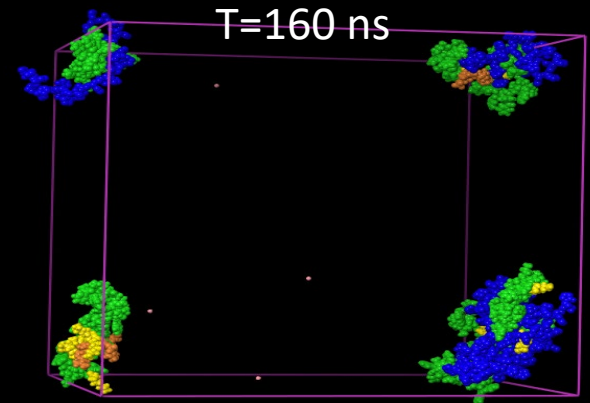
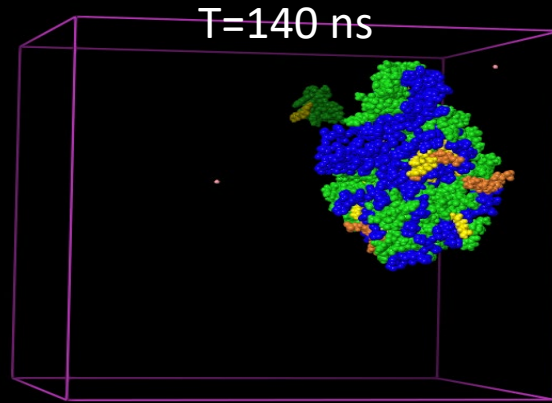
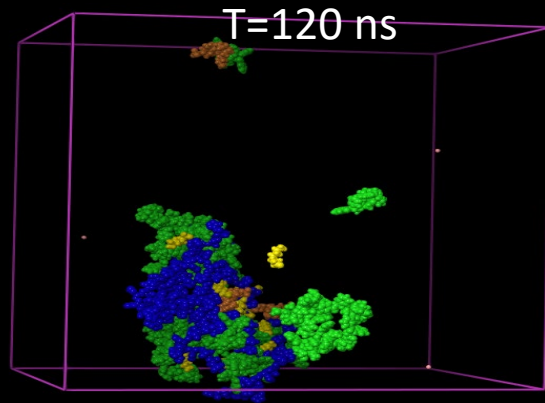
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Nanoparticle simulation snapshots...



Nanoparticle simulation snapshots...



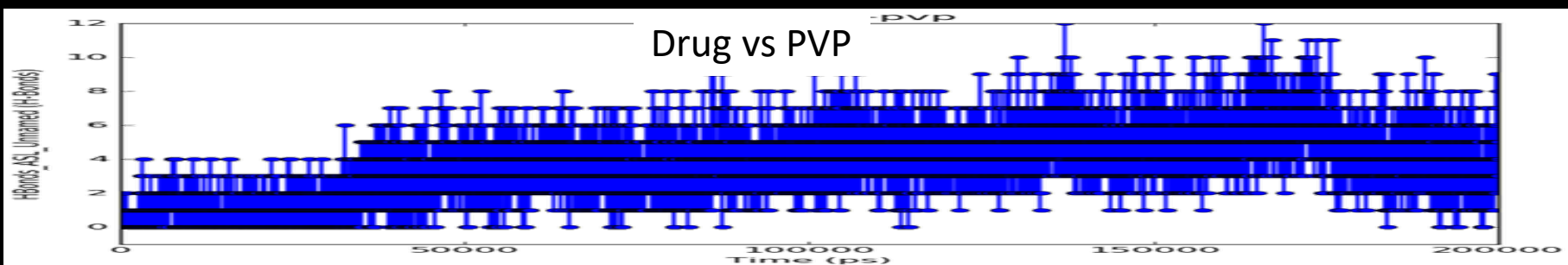
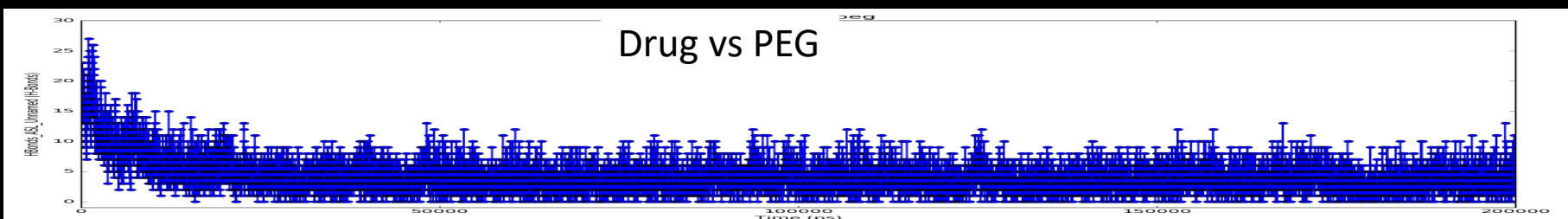
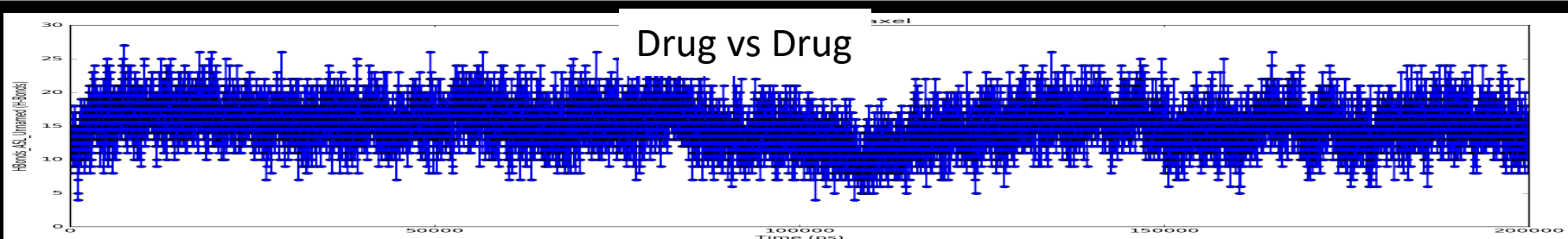
Analysis...



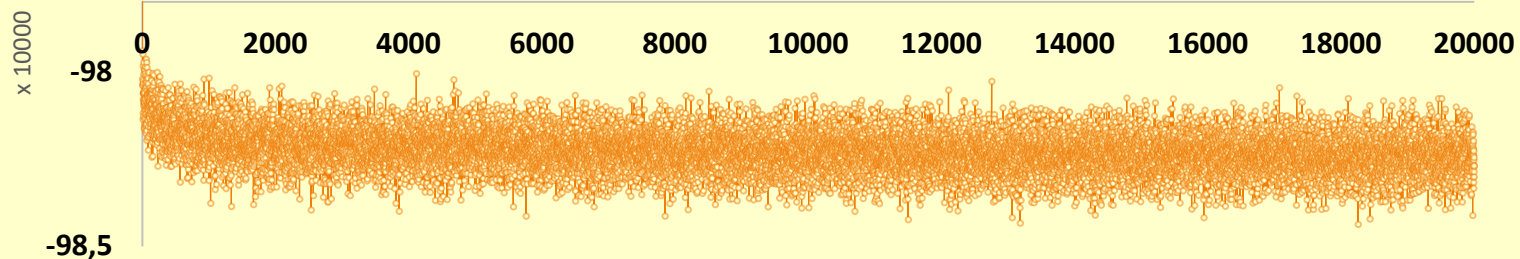
H-bond analysis

RMSD

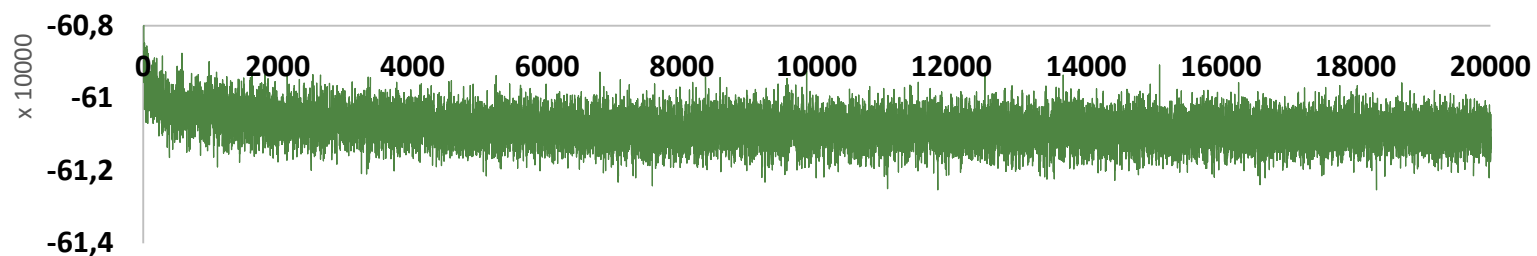
Potential energy



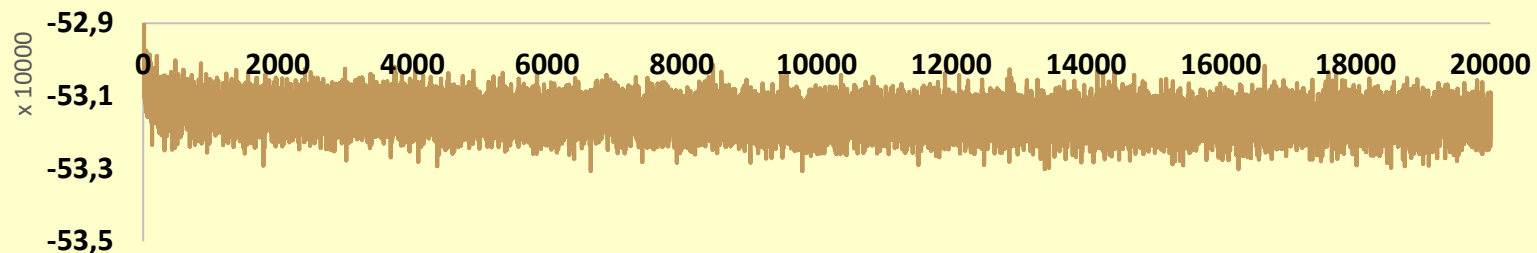
Potential Energy vs Time



Full system



System -PVP



System -PEG

Run Time: 200ns

Ensemble: NPT

Energies: Kcal/mol

Other systems studied...

System	Drug	NoPVP	NoPEG	NoFA	NoCHO
Average_Drug-Drug	15.14	31.28	32.80	23.57	29.58
Average_Drug-PVP	3.45	-	7.93	9.24	8.18
Average_Drug-PEG	4.04	15.23	-	16.10	14.91
Average_Drug-FA	0.48	4.28	5.84	-	2.85
Average_Drug-CHO	0.053	0.75	1.36	0.086	-

Removal of PVP, PEG, FA and CHO increases Drug-Drug H-bonds

Role of PEG seems increasing as soon as any of the excipients are removed – which means that in the absence of any of the components Drug can still form aggregates in the presence of PEG

- ❑ Composition is as per the experimental stoichiometry
- ❑ Simulation carried for 200 ns – parameters studies are stable
- ❑ Studies conducted also without PVP, PEG, CHO, FA. PVP and PEG has a larger role in NP stabilization in comparison with CHO and FA
- ❑ Length of the nanoparticle ~10nm (exclusion – PEG, water, EtOH)
- ❑ Width is ~7nm
- ❑ This study can help us understand the role of each excipient in such formulations and help in optimizing formulations
- ❑ Computational optimization can be leveraged for difficult to formulate drugs

Acknowledgments



Dr. C. T. Rao
Head Drug Discovery

Dr. Ajay Khopade
VP, FDD

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And

Sun Pharma Advanced Research Company Limited
Vadodara, Gujarat, India



Thank You

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