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## 1 A 6+T Parametrization

- 2 The algorithm followed for the distance-based loss function optimization is detailed below.

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**Algorithm 1** 6+T parametrization used in UMD-fit

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1: optim = LBFGS(params=[euler, trans, torsions], lr=0.5)
2: for i in range(max_steps):
3:     def closure():
4:         optim.zero_grad()
5:         coords = coords + trans
6:         rot_mat = rot_from_euler(euler)
7:         coords = einsum('rc,nc->nr', rot_mat, coords - com) + com
8:         aux_coords = coords.clone()
9:         for (i,j,k,l), value in zip(torsion_idxs, torsion_values):
10:             coords = update_dihedral(coords, idxs=[i,j,k,l], value=value)
11:             coords = kabsch(coords, aux_coords)
12:             loss = loss_func(coords, pocket_coords, pred_dist_mat)
13:             loss.backward()
14:     return loss
15: loss = optimizer.step(closure)
```

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- 3 More details on the performance of  $6 + T$  alone and the stereochemistry preservation in UMD-fit  
4 relative to Uni-Mol Docking are shown in Table 1. The difference can be largely attributed to higher  
5 quality conformers in the distance optimization post-processing, but not improved modelling by the  
6 ligand processing module in Uni-Mol. We hypothesize this might be explained by incorrect training  
7 data (wrong stereochemistry) in original Uni-Mol training, thus yielding a model less sensitive to  
8 those features and specific geometries. Further work might involve retraining the original model with  
9 correct input features.

Table 1: Performance for different inference strategies for Uni-Mol Docking in the CASF-2016 test set

Strategy	RMSD ( $\downarrow$ , Å)	symRMSD ( $\downarrow$ , Å)	% $\leq$ symRMSD ( $\uparrow$ )					
			0.5Å	1.0Å	1.5Å	2.0Å	3.0Å	5.0Å
Baseline	1.67	1.54	14.74	56.14	72.98	81.05	87.72	94.39
6+T	1.81	1.65	10.18	45.96	70.53	79.29	85.61	93.68
UMD-fit	1.75	1.57	12.63	51.58	72.28	80.70	87.72	94.03