

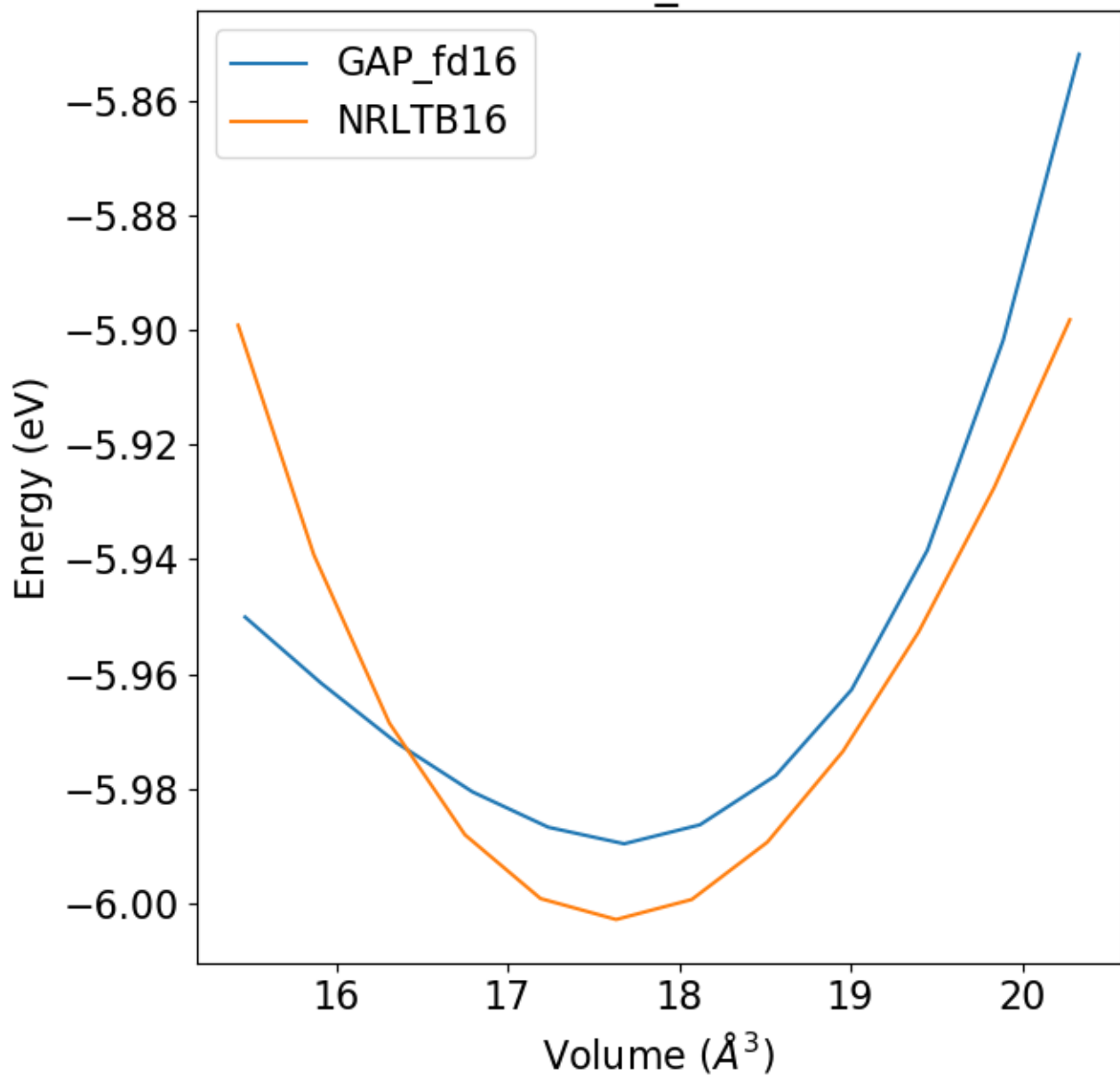
GAP_fd16

```
<command_line><![CDATA[ at_file=Ti_hcp_bcc16.xyz gap="distance_Nb order=2 cutoff=9.0 covariance_type=ard_se n_sparse=20
theta_uniform=1.0 delta=0.003 compact_clusters=T :soap cutoff=4.0 covariance_type=dot_product n_sparse=200 delta=0.095
sparse_method=cur_points l_max=10 n_max=10 zeta=4 atom_sigma=0.5" e0=-4.334618118602442 sparse_separate_file=F
gp_file=gp_soap_hcp_bcc_fdcell.xml default_sigma={0.0001 0.05 0.05 0.0}]]></command_line>
```

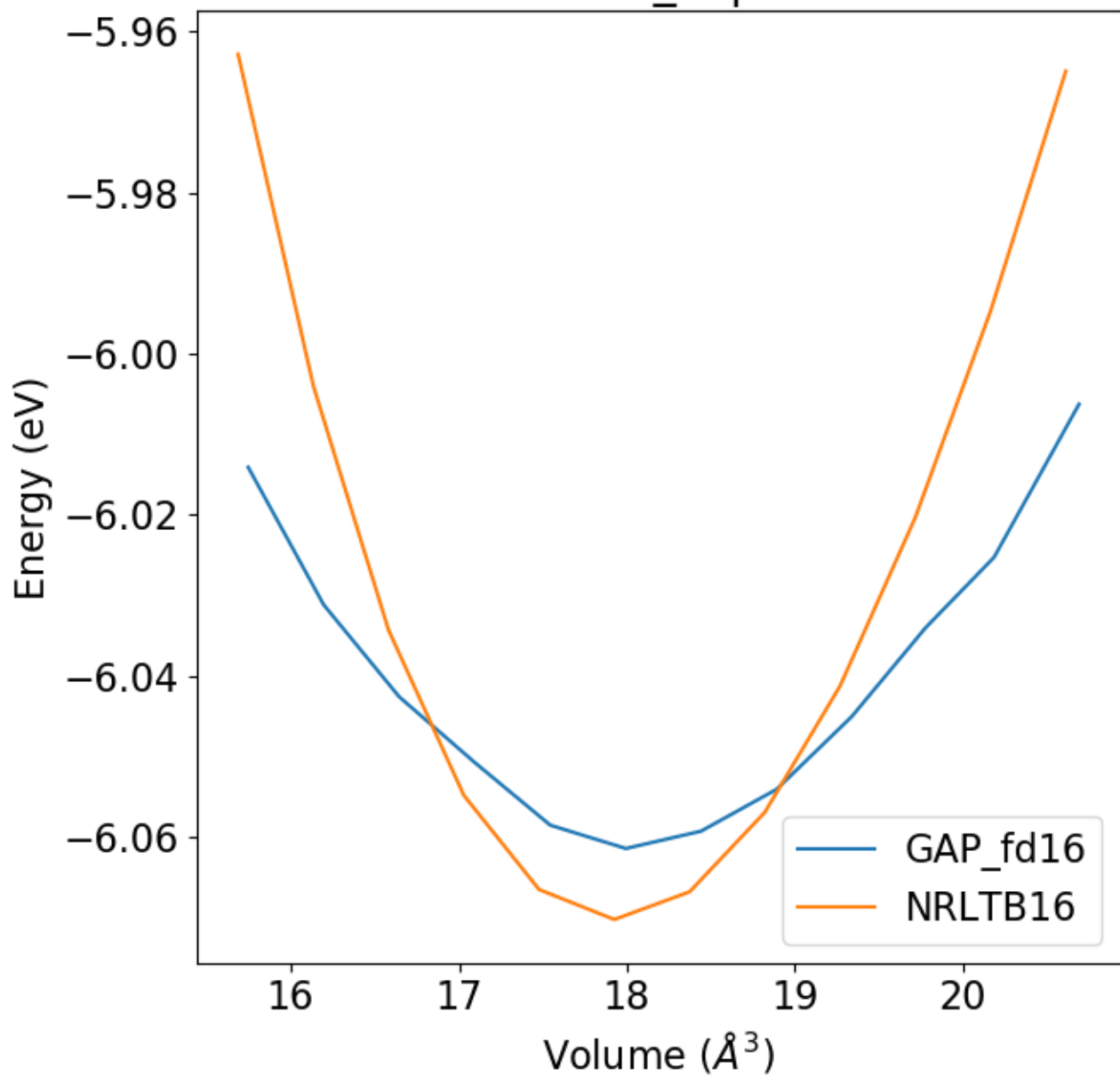
{'hcp_fdcell': 216, 'bcc_fdcell': 162}

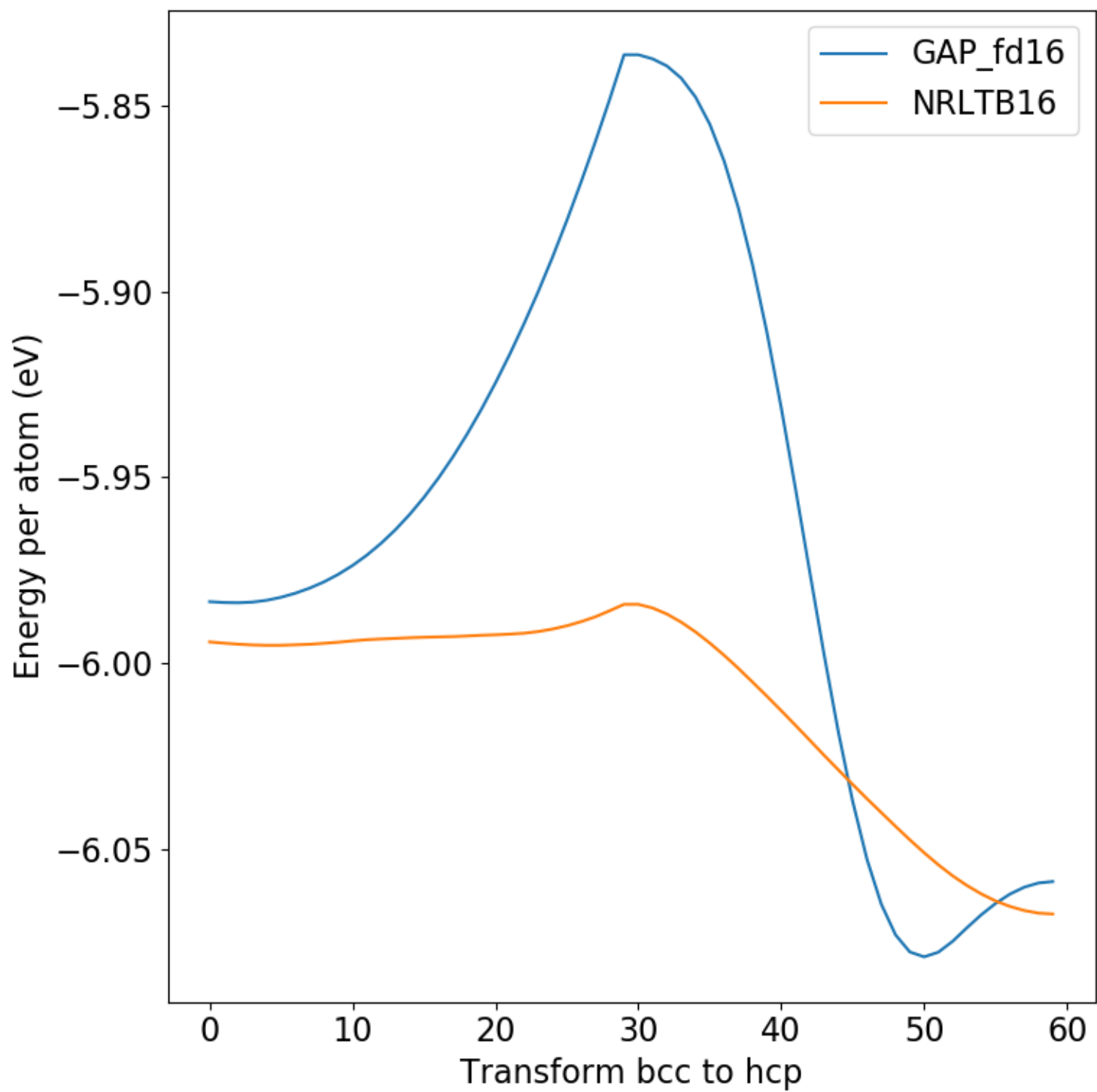
-		NRLTB16	GAP_fd16
0	bcc_c12	100.20	-44%
1	bcc_c11	105.31	-3%
2	bcc_c44	38.92	192%
3	hcp_c44	34.28	140%
4	hcp_c13	68.39	-2%
5	hcp_c12	57.67	-151%
6	hcp_c11	176.57	-6%
7	hcp_c66	56.05	86%
8	hcp_c33	171.23	56%

bulk_bcc



bulk_hcp





dimer

