

1 Command Line Dictionary

acf	yes no{no}
amsklp	automask low-pass limit(in Å)
angerr	angular error(in degrees)
append	yes no{no}
astigerr	astigmatism error(in degrees)
avg	yes no
bfac	bfactor
bfacerr	bfactor error
bin	binarize{no}
box	image size(in pixels)
boxtab	boxfiles.txt
center	yes no{no}
chunksz	chunk size
class	cluster2extract
clip	clip2box{256}
clustvalid	yes homo no {no}
comlindoc	shc_clustering_nclsX.txt
compare	yes no{no}
corner	corner size(in pixels)
countvox	yes no{no}
cs	spherical aberration constant(in mm){2.7}
ctf	yes no flip mul{no}
ctfsq	yes no{no}
ctfstats	yes no{no}
cube	cube side(in pixels)
deferr	defocus error(in microns)
defocus	defocus(in microns)
deftab	text file with CTF info
dens	density
dir	directory
discrete	yes no{no}
diversify	yes no{yes}
doclist	list of oritabs
dynlp	yes no{yes}
e1	1st Euler angle
e2	2nd Euler angle
e3	3d Euler angle
edge	edge size(in pixels)
eo	yes no{yes}
errify	yes no{no}
even	yes no{no}
fbody	body of file
filetab	movies.txt
find	Fourier index
fname	filename
frac	fraction
frac	fraction of amplitude contrast{0.07}
fromp	from particle index

fsc	fsc_state1.bin
ft2img	yes no{no}
guinier	yes no{no}
hfun	sigm tanh lin{sigm}
hist	var2plot
hp	high-pass limit(in Å)
inner	inner mask radius(in pixels)
kv	acceleration voltage{300}
label	class state subclass{class}
lp	low-pass limit(in Å){20}
lpstart	starting low-pass limit
lpstop	stay at this low-pass limit
masscen	yes no{no}
maxits	maximum number of iterations{500}
minp	minimum cluster population
mirr	no 2d 3d{no}
moldiam	molecular diameter(in Å)
msk	mask radius(in pixels)
mskfile	mask.ext
msktype	hard soft{soft}
mul	shift multiplication factor
mw	molecular weight(in kDa)
ncls	nr of clusters
ndiscrete	nr of discrete oris
ndocs	nr of documents
neg	yes no
newbox	new box size(in pixles)
nframes	nr of movie frames
noise	yes no{no}
noise_norm	yes no
norm	yes no{no}
npart	nr of partitions(nodes)
npeaks	nr of correlation peaks
nptcls	nr of particles
nran	size of random sample
nspace	nr of orientations in discrete space
nstates	nr of conformational states
nthr	nr of threads (CPUs within a socket)
nvox	nr of voxels
odd	yes no{no}
oritab	orientations
oritab2	orientations
outfile	output text file
outstk	output image stack
outvol	outvol.ext
pgrp	cn dn t o i{c1}
phrand	yes no{no}
plot	yes no{no}
pspecsz	box size of power spectrum
refine	no shc{no}

refs	initial_references.ext
rnd	yes no{no}
round	yes no{no}
scale	scale factor
shalgn	yes no{no}
sherr	shift error(in pixels)
smpd	sampling distance(in Å)
snr	signal2noise ratio
soften	yes no{no}
split	nr of partitions to split into
srch_inpl	yes no{yes}
startit	start iteration nr
state	state index
stats	yes no print{no}
stk	input particle stack
stk2	input particle stack nr 2
stk3	input particle stack nr 3
thres	threshold
top	stop particle index
trs	origin shift range[-trs,trs](in pixels)
vis	yes no{no}
vol1	invol.ext
vol2	invol2.ext
which_iter	iteration number
width	pixels width
xdim	nr of pixels in x-dim
xsh	x shift(in pixels)
ydim	nr of pixels in y-dim
ysh	y shift(in pixels)
zero	yes no{no}
zsh	z shift(in pixels)