

Traceback (most recent call last):  
  File "E:\ChimeraX 1.8\bin\Lib\site-packages\chimerax\open\_command\dialog.py", line 334, in \_qt\_safe  
    run(session, "open " + " ".join([FileNameArg.unparse(p) for p in paths]) + (""  
  File "E:\ChimeraX 1.8\bin\Lib\site-packages\chimerax\core\commands\run.py", line 49, in run  
    results = command.run(text, log=log, return\_json=return\_json)  
              ^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^  
  File "E:\ChimeraX 1.8\bin\Lib\site-packages\chimerax\core\commands\cli.py", line 3213, in run  
    result = ci.function(session, \*\*kw\_args)  
             ^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^  
  File "E:\ChimeraX 1.8\bin\Lib\site-packages\chimerax\open\_command\cmd.py", line 131, in cmd\_open  
    models = Command(session, registry=registry).run(provider\_cmd\_text, log=log)[0]  
             ^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^  
  File "E:\ChimeraX 1.8\bin\Lib\site-packages\chimerax\core\commands\cli.py", line 3213, in run  
    result = ci.function(session, \*\*kw\_args)  
             ^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^  
  File "E:\ChimeraX 1.8\bin\Lib\site-packages\chimerax\open\_command\cmd.py", line 213, in provider\_open  
    models, status = collated\_open(session, None, [data], data\_format, \_add\_models,  
                     ^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^^  
  File "E:\ChimeraX 1.8\bin\Lib\site-packages\chimerax\open\_command\cmd.py", line 513, in collated\_open  
    return remember\_data\_format()  
           ^^^^^^^^^^^^^^^^^^^^^^  
  File "E:\ChimeraX 1.8\bin\Lib\site-packages\chimerax\open\_command\cmd.py", line 484, in remember\_data\_format  
    models, status = func(\*func\_args, \*\*func\_kw)  
                     ^^^^^^^^^^^^^^^^^^^^^^^^^^^  
  File "C:\Users\Administrator\AppData\Local\UCSF\ChimeraX\1.8\Python311\site-packages\SEQCROW\\_\_init\_\_.py", line 596, in open  
    return open\_aarontools(  
           ^^^^^^^^^^^^^^^^  
  File "C:\Users\Administrator\AppData\Local\UCSF\ChimeraX\1.8\Python311\site-packages\SEQCROW\io.py", line 45, in open\_aarontools  
    raise e  
  File "C:\Users\Administrator\AppData\Local\UCSF\ChimeraX\1.8\Python311\site-packages\SEQCROW\io.py", line 33, in open\_aarontools  
    fr = FileReader(  
         ^^^^^^^^^^^  
  File "C:\Users\Administrator\AppData\Local\UCSF\ChimeraX\1.8\Python311\site-packages\AaronTools\fileIO.py", line 1555, in \_\_init\_\_  
    self.read\_orca\_out(f, get\_all, just\_geom)  
  File "C:\Users\Administrator\AppData\Local\UCSF\ChimeraX\1.8\Python311\site-packages\AaronTools\fileIO.py", line 2568, in read\_orca\_out  
    \_, \_, rot = line.split()  
    ^^^^^^^^^  
ValueError: not enough values to unpack (expected 3, got 0)  
  
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File "C:\Users\Administrator\AppData\Local\UCSF\ChimeraX\1.8\Python311\site-packages\AaronTools\fileIO.py", line 2568, in read\_orca\_out  
\_, \_, rot = line.split()  
^^^^^^^^^  
  
*See log for complete Python traceback.*