Say I had a list of ligands that were ranked according to their binding energy from a docking program. The output from the docking program had ligand numbers that corresponded with the .sdf file I exported from DataWarrior after converting to conformers. Would it be possible to take this list of row numbers and use the row list function to view these ligands in order of ranked binding energy? I'm not sure if row lists can be used this way in Data Warrior. I have not tried it yet.