**Instructions: Bocso’s class II motif**

1. Create a csv table with the following headers:
2. Create a folder containing this csv file, the motif.py and the 2 clustalw files
3. Right-click on motif.py and run with python launcher
4. Important: If you double-click motif.py, it might be that it only opens in a text editor
5. This creates a fasta file called sorted\_and\_aligned\_peptides
6. Sri recommended to use Meme motif finder (<http://meme-suite.org/tools/meme>)
7. Copy-paste sequences, select “one occurrence per sequence” and press go