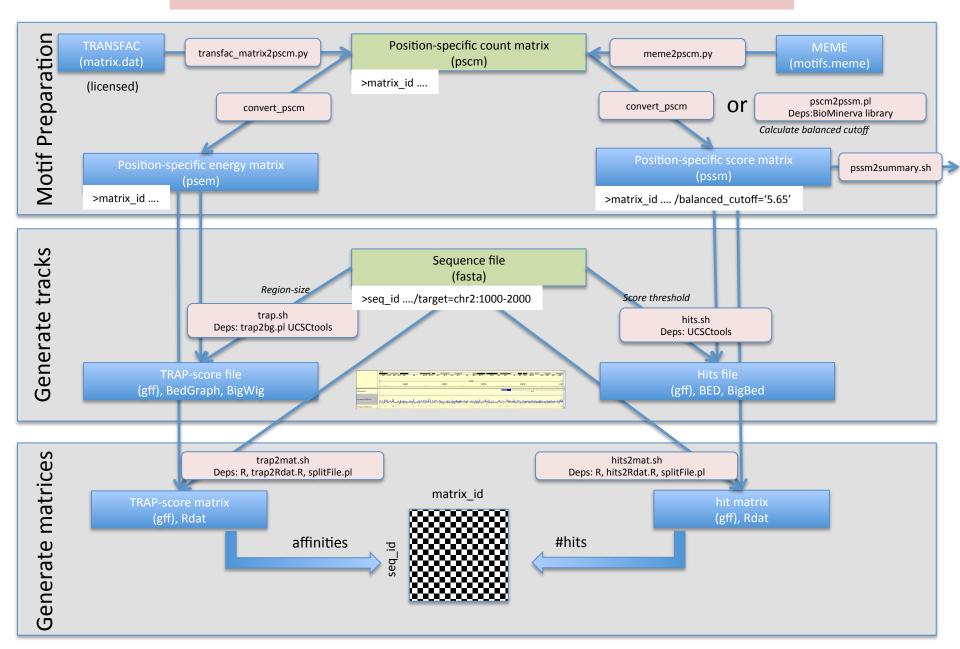
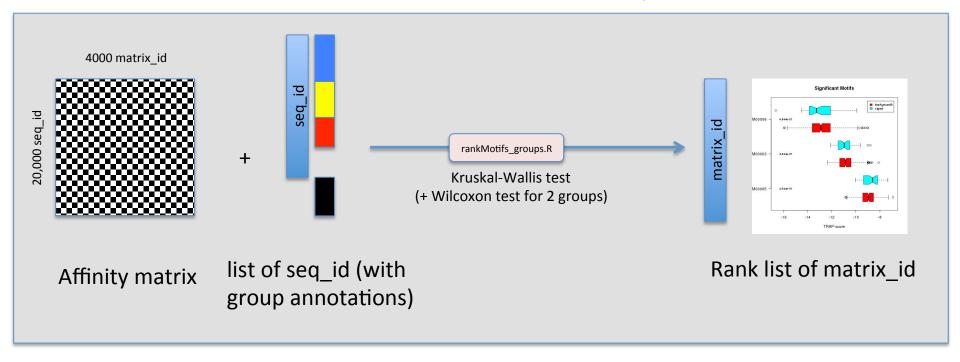
TRANSFAC UTILITIES



Contributing Developers: Morgane Thomas-Chollier, Szymon Kielbasa, Helge Roider, Sarah Diehl, Abinhab Ray, Thomas Manke

Downstream Analysis



Notice that for quantitative assignment of seq_ids to a single group (e.g. tissues) an iterative association test has also been developed:

Roider HG1, Manke T, O'Keeffe S, Vingron M, Haas SA. PASTAA: identifying transcription factors associated with sets of co-regulated genes. Bioinformatics. 2009 Feb 15;25(4):435-42. doi: 10.1093/bioinformatics/btn627. Epub 2008 Dec 9.