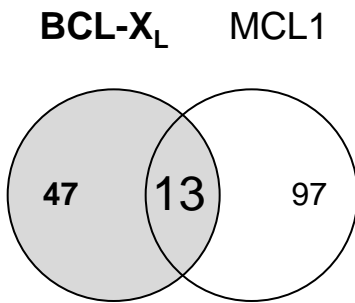
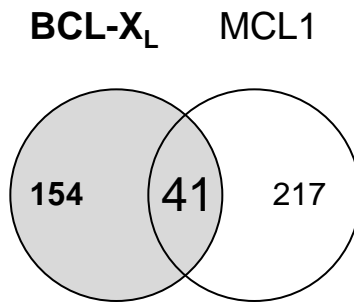


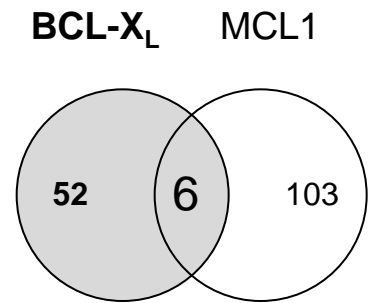
Computational predictions



TargetScan 5.1
(<http://www.targetscan.org/>)



miRanda
(<http://www.microrna.org/>)



DIANA - microT v3.0
(<http://diana.cslab.ece.ntua.gr/>)



11 putative miRNAs

(intersection between at least two algorithms)

hsa-miR-:

17-5p, 32, 96, 133a, 133b, 182,
193b, 204, 214, **491-5p**, 504