



SecretomeP 2.0a Server - prediction results

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Non-classically secreted proteins should obtain an NN-score / SecP score exceeding the threshold, but not at the same time be predicted to contain a signal peptide.

The recommended thresholds are **0.5** for bacterial sequences and **0.6** for mammalian sequences.

Network 1	Network 2	Network 3	SecP score	Sequence name
0.065498	0.107936	0.137881	0.103772	AP15_new2331_APP8_00109_1331
0.086907	0.861762	0.191545	0.380071	AP15_new2331_APP8_00138_1305
0.059357	0.039810	0.115986	0.071718	AP15_new2331_APP8_00178_1270
0.068116	0.008114	0.103679	0.059970	AP15_new2331_APP8_00203_1246
0.929563	0.999947	0.878040	0.935850	AP15_new2331_APP8_00382_1155
0.053000	0.129206	0.124553	0.102253	AP15_new2331_APP8_00717_959
0.058856	0.326073	0.103123	0.162684	AP15_new2331_APP8_00762_924
0.074055	0.207839	0.189233	0.157042	AP15_new2331_APP8_00885_842
0.050498	0.037906	0.104238	0.064214	AP15_new2331_APP8_00890_838
0.347511	0.064949	0.386512	0.266324	AP15_new2331_APP8_00933_1620
0.034456	0.005692	0.113247	0.051132	AP15_new2331_APP8_01026_754
0.230942	0.038236	0.272693	0.180624	AP15_new2331_APP8_01031_759
0.106500	0.012407	0.158691	0.092533	AP15_new2331_APP8_01035_763
0.259802	0.104518	0.385090	0.249803	AP15_new2331_APP8_01069_791
0.931694	0.999763	0.879001	0.936819	AP15_new2331_APP8_01078_1723
0.243977	0.405886	0.185579	0.278481	AP15_new2331_APP8_01385_1549
0.045957	0.065131	0.084324	0.065137	AP15_new2331_APP8_01477_1563
0.868756	0.966625	0.788182	0.874521	AP15_new2331_APP8_01587_696
0.144056	0.057704	0.174509	0.125423	AP15_new2331_APP8_01753_407
0.053302	0.021478	0.073441	0.049407	AP15_new2331_APP8_01963_253
0.042290	0.300273	0.166034	0.169532	AP15_new2331_APP8_01976_240
0.059693	0.025756	0.072426	0.052625	AP15_new2331_APP8_02011_472
0.047426	0.085255	0.129206	0.087296	AP15_new2331_APP8_02015_468
0.082262	0.004992	0.090792	0.059349	AP15_new2331_APP8_02063_1372
0.044916	0.705370	0.119413	0.289900	AP15_new2331_APP8_02182_553

Explain the output. [Go back](#).