

# Result of Medical Clustering task of ImageCLEF 2015 at CLEF2015

Name	Name of the Sumited file	Exact Match	Any Match	Hamming similary
IBM MMAFL	1431923404568__IBM_MRL_ef_mslbpyr_scodes_densemblbp_densesift_th.txt	<b>0.752</b>	<b>0.864</b>	0.863
SNUMedInfo	1431946260816__SNUMedinfo3.txt	0.709	0.856	0.895
SNUMedInfo	1431946236163__SNUMedinfo2.txt	0.699	0.844	0.890
IBM MMAFL	1431923215231__IBM_MRL_ef_mslbpyr_scodes_densemblbp_densesift_max.txt	0.695	0.840	0.889
IBM MMAFL	1431773613950__IBM_MRL_Sift_FV.txt	0.692	0.832	<b>0.896</b>
IBM MMAFL	1431773914574__IBM_MRL_SMO_edge_histogram_pyramid23.txt	0.692	0.732	0.755
IBM MMAFL	1431932028897__fuse_pred.txt	0.689	0.820	0.890
SNUMedInfo	1431946207517__SNUMedinfo1.txt	0.679	0.820	0.879
IBM MMAFL	1431922644645__IBM_MRL_mlhist_mslbp_pyr_max.txt	0.672	0.812	0.874
AmrZEGY	1431858638236__run1.txt	0.646	0.780	0.868
NLM	1431352031992__OpenI_SetA_5class.out	0.613	0.740	0.849
CASMIP	1431623161365__C_knnclassification.txt	0.606	0.732	0.843
CASMIP	1431621989456__svmclassification.txt	0.603	0.728	0.847
IBM MMAFL	1431773973109__IBM_MRL_SMO_lbp_histogram_pyramid23.txt	0.603	0.708	0.778
CASMIP	1431399984301__knnclassification.txt	0.599	0.724	0.847
IBM MMAFL	1431922924104__IBM_MRL_scodes_denseMBLBP_max.txt	0.599	0.724	0.838
NLM	1431634802808__OpenI_5class.out	0.593	0.716	0.842
CASMIP	1431623320464__TC_knnclassification.txt	0.589	0.712	0.842
CASMIP	1431623385130__TC_logclassification.txt	0.589	0.712	0.842
CASMIP	1431623447485__TC_logclassification.txt	0.589	0.712	0.842
CASMIP	1431622163959__logclassification.txt	0.573	0.692	0.833
CASMIP	1431622397898__dbnclassification.txt	0.573	0.692	0.833
CASMIP	1431623254773__TC_dbnclassification.txt	0.563	0.680	0.830
CASMIP	1431622528178__T_knnclassification.txt	0.550	0.664	0.826
NLM	1431635462767__SetA_5class.out	0.543	0.656	0.810
IBM MMAFL	1431773776203__IBM_MRL_DF_lbp_histogram_pyramid23.txt	0.510	0.616	0.849
BMET	1431903128194__clustering_run_1.txt	0.497	0.596	0.816
IBM MMAFL	1431773817715__IBM_MRL_DF_edge_histogram_pyramid23.txt	0.470	0.568	0.835
db Lab	1431790361993__results.txt	0.219	0.264	0.664

Exact Match: we counted correct if at a bit location ground truth says 1 and your system says 1, also how many of the true negatives your system has correctly identified

Any Match: we counted a correct classification if your system says 1 and at that location ground truth is also 1, also how many of the true negatives your system has correctly identified

Hamming Similarity: calculation of hamming distance for of your output and ground truth, not a good measure because of too many zeros