	queryid	subjectids	identity	alignmentl	mismatche	gapopens	q.start	q.end	s.start	s.end	evalue	bitscore	positives	mlog.evalu	pdb.id	acc
1	Query_567	4IZ6_A	80	5	1	0	3	7	379	383	7.9	21.4	100	-2.06686	4IZ6_A	4IZ6_A
2	Query_567	3RG2_A	80	5	1	0	3	7	361	365	7.9	21.4	100	-2.06686	3RG2_A	3RG2_A
3	Query_567	6IYK_A	80	5	1	0	3	7	359	363	7.9	21.4	100	-2.06686	6IYK_A	6IYK_A