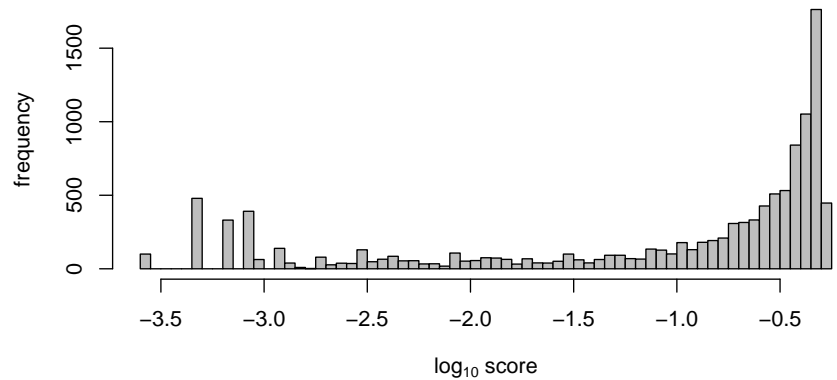


Report for PSM Secion in mzTab File example

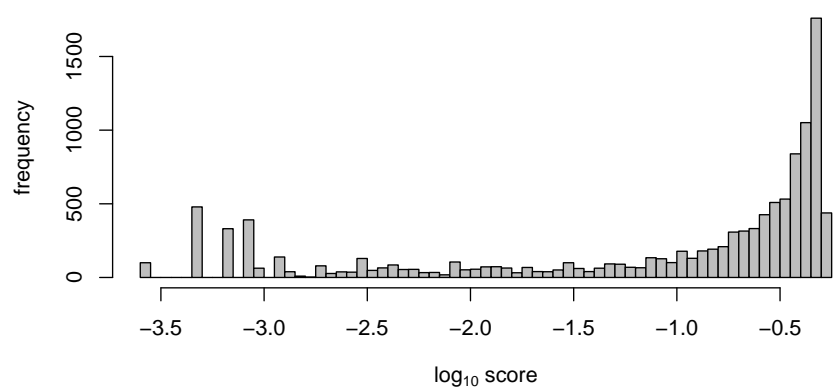
The analysis identified a total of 15,001 PSMs.

	total	non-redundant
PSMs	15,001	9,789
unique PSMs	6,890	4,758
target PSMs	14,978	9,770
decoy PSMs	0	0
target + decoy PSMs	23	19

Table 1: Number of PSMs. Two PSMs are considered *non-redundant* if they differ in amino acid sequence. They are considered *redundant*, if they differ merely in modifications. A PSM is considered *unique*, if its amino acid sequence can be mapped uniquely to a single protein. The sequence can appear in either target proteins, decoy proteins or both of them.

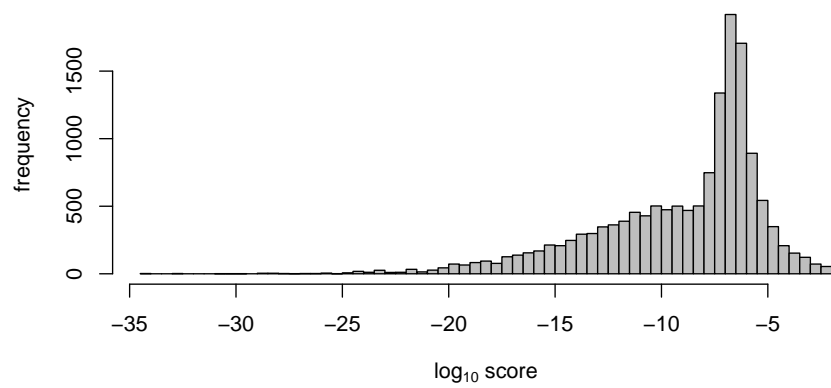


(a) 15,001 PSMs

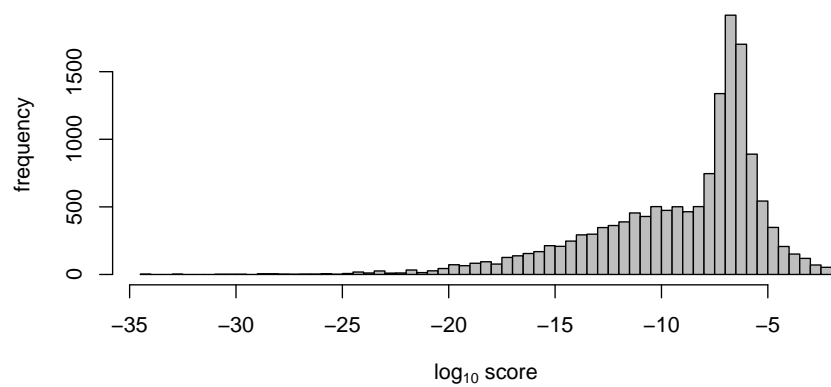


(b) 14,978 target PSMs

Figure 1: Score distribution of `search_engine_score[1]`.



(a) 15,001 PSMs



(b) 14,978 target PSMs

Figure 2: Score distribution of e-values.