

Sample	Redundancy	UniProt	Sequences in search space	Identified protein groups	Identified peptides	Identified PSMs
HCT116	No	/	92 931	4 322	28 729	179 175
	Yes	/	133 051	4 330	28 684	179 056
Jurkat	Yes	/	201 419	4 450	28 689	172 273
HCT116	Yes	Canonical	176 202	4 333	28 402	177 473
		Spliced	186 627	4 347	28 372	176 978
Jurkat	Yes	Spliced	253 734	4 477	28 548	171 116
HCT116, only UniProt		Canonical	71 356	4 294	28 443	180 526