

reads against *Mycobacterium smegmatis* and *Mycobacterium gilvum* genomes.
Each entry contains a pair “Fraction (in classified closer to *M.smegmatis*
/ Fraction of reads classified closer to *M.gilvum*”.

	weight				
	14	16	18	20	22
contig hit nb	52/41	39/48	24/37	11/24	07/17
contig cover	54/42	44/47	25/37	11/24	06/17
contig Jaccard	30/70	35/61	23/42	11/26	06/18
spaced hit nb	51/40	34/47	20/40	12/32	08/23
spaced cover	53/42	39/51	21/42	12/32	08/25
spaced Jaccard	28/72	32/66	20/50	11/33	08/27