

Accession	Description	Score	Coverage	# Proteins	# Unique	# Peptides	% of Unique	% of Peptide	% of Unique	% of Peptide	# PSMs	# AAs	MW [kDa]	calc. pI	% of Unique Peptides/[kDa] (Total % of Peptides/[kDa] (Total))	
A0A216P6P2	Major capsid protein OS=Salmonella phage BPS1556 OX=2060121 GN=BPS1556_11 PE=3 SV=1 - [A0A216P6P2_9CAUD]	570,155441	58,15	1	4	21	0,85470085	2,27027027	0,02059132	0,054695	199	368	41,5078184	5,74462891	1,864508166	1,705051225
A0A251GPP5	Putative structural protein OS=Enterobacteria phage vB_EcoM_IME338 OX=2163888 PE=4 SV=1 - [A0A251GPP5_9CAUD]	461,879526	55,71	9	17	32	6,3247863	3,45945946	0,06294029	0,05994327	141	508	57,130863	6,37939453		
A0A689W2I9	Tail tapemeasure protein OS=Escherichia phage pinkflif OX=2696440 GN=pinkflif_21 PE=4 SV=1 - [A0A689W2I9_9CAUD]	390,625679	45,24	1	3	31	0,85470085	3,35135135	0,01065481	0,04177839	105	736	80,2173498	9,67138672		
A0A455XBP6	phage tail tape measure OS=Escherichia phage L27 OX=2562890 PE=4 SV=1 - [A0A455XBP6_9CAUD]	380,828899	54,42	1	3	39	0,64102564	4,21621622	0,0079953	0,05258747	109	735	80,157295	9,65673828		
A0A0H3UC50	Major capsid protein OS=Escherichia phage HY02 OX=1527531 GN=HY02_058 PE=3 SV=1 - [A0A0H3UC50_9CAUD]	375,196736	58,15	1	1	19	0,21367521	2,05405405	0,00513717	0,04938346	113	368	41,5999644	5,77001953		
A0A689XCT5	Major capsid protein OS=Escherichia phage warpig OX=2696461 GN=warpig_95 PE=3 SV=1 - [A0A689XCT5_9CAUD]	375,196736	63,86	1	1	19	0,21367521	2,05405405	0,00515279	0,04935365	113	368	41,4678487	5,63037109		
A0A481W3F5	Tail length tape-measure protein OS=Escherichia phage PH811 OX=2530036 PE=4 SV=1 - [A0A481W3F5_9CAUD]	362,587187	55,1	1	3	39	0,64102564	4,21621622	0,00798564	0,05252394	105	735	80,2727219	9,67138672		
A0A455XHL7	phage protein OS=Escherichia phage L27 OX=2562890 PE=4 SV=1 - [A0A455XHL7_9CAUD]	322,148818	66,87	2	2	23	0,42735043	2,48648649	0,00805727	0,04688204	97	489	53,0391207	4,66552734		
A0A689WQ1	Baseplate assembly protein OS=Escherichia phage skuden OX=2696443 GN=skuden_43 PE=4 SV=1 - [A0A689WQ14_9CAUD]	288,274184	66,46	2	3	23	0,64102564	2,48648649	0,01207721	0,04682905	86	489	53,097898	4,61474609		
A0A3845J16	Tail fibers protein OS=Enterobacteria phage XTG1 OX=1651201 PE=4 SV=1 - [A0A3845J16_9CAUD]	286,823802	29,97	1	1	18	0,21367521	1,94594595	0,00248917	0,02266898	91	794	85,8417953	8,77783203		
A0A689XFQ2	Putative tail protein OS=Escherichia phage radamba OX=2696441 GN=radamba_56 PE=4 SV=1 - [A0A689XFQ2_9CAUD]	282,961296	44,71	1	12	12	2,56410256	1,29729793	0,08181677	0,04139487	76	293	31,3395703	5,02099609		
A0A3845J98	Tail fibers protein OS=Enterobacteria phage KhF2 OX=1651204 PE=4 SV=1 - [A0A3845J98_9CAUD]	266,785674	26,39	1	2	18	0,42735043	1,94594595	0,00486625	0,02270497	86	792	85,7057226	8,49951172		
A0A689WPE1	Tail fibers protein OS=Escherichia phage skuden OX=2696443 GN=skuden_39 PE=4 SV=1 - [A0A689WPE2_9CAUD]	257,581438	24,33	1	3	18	0,64102564	1,94594595	0,00765295	0,02323186	88	781	83,7619343	8,58740234		
A0A251GP51	Tail fiber protein OS=Enterobacteria phage vB_EcoM_IME338 OX=2163888 PE=4 SV=1 - [A0A251GP51_9CAUD]	240,493549	22,86	1	4	18	0,85470085	1,94594595	0,01003028	0,02283652	78	783	85,2120239	6,87353516		
A0A384FM4	Major capsid protein (Fragment) OS=Escherichia phage VNV OX=2488710 PE=4 SV=1 - [A0A384FM4_9VIRU]	214,377004	54,17	1	2	8	0,42735043	0,86486486	0,02664768	0,05392915	50	144	16,0370566	7,59130859		
A0A689M307	Tail fiber protein OS=Shigella phage Z31 OX=2675202 PE=4 SV=1 - [A0A689M307_9CAUD]	198,247812	25,79	1	2	18	0,42735043	1,94594595	0,00497566	0,02265675	70	795	85,8881497	8,86572266		
A0A482MR91	Tail fiber OS=Escherichia phage vB_EcoM_LMP33 OX=2491664 PE=4 SV=1 - [A0A482MR91_9CAUD]	179,12171	20,78	2	2	16	0,42735043	1,72972973	0,00493533	0,01997607	63	794	86,95900727	8,04541016		
A0A482HJG1	Uncharacterized protein OS=Shigella phage Silverhawkium OX=2530185 GN=Silverhawkium_gp81 PE=4 SV=1 - [A0A482HJG1_9CAUD]	164,812734	29,1	1	16	16	0,21367521	1,72972973	0,00385681	0,03121236	52	488	55,4021215	6,31591797		
A0A384T024	Tail fiber protein OS=Enterobacteria phage KhF3 OX=1651200 PE=4 SV=1 - [A0A384T024_9CAUD]	150,392019	41,27	3	10	10	2,13675214	1,08108108	0,05100214	0,02580432	54	395	41,8953453	6,96142578		
A0A482MRM	Uncharacterized protein OS=Escherichia phage vB_EcoM_LMP25 OX=2491663 PE=4 SV=1 - [A0A482MRM9_9CAUD]	125,443554	59,04	3	8	8	1,70940171	0,86486486	0,09384935	0,0474827	37	166	18,2143167	18,14794922		
A0A45XN56	Uncharacterized protein OS=Salmonella phage DaR_2019a OX=2571168 GN=FFZ30_0023 PE=4 SV=1 - [A0A45XN56_9CAUD]	117,398192	49,82	8	13	13	2,77777778	1,40540541	0,08778159	0,04441274	36	285	31,6441945	5,79541016		
A0A689SPQ7	Uncharacterized protein OS=Escherichia phage JN01 OX=2692737 PE=4 SV=1 - [A0A689SPQ7_9CAUD]	110,11253	72	1	6	9	1,28205128	0,97297297	0,0940883	0,07140539	39	125	13,6260432	9,94775391		
A0A689LU8	Uncharacterized protein OS=sphage NBeCO05 OX=2712974 PE=4 SV=1 - [A0A689LU8_9CAUD]	101,938223	48,22	1	4	20	0,85470085	2,16216216	0,014922	0,04425054	31	450	48,8618259	5,18603516		
A0A0E3TRX4	Uncharacterized protein OS=Escherichia phage vB_EcoM_AYO145A OX=1636202 GN=AYO145A_064 PE=4 SV=1 - [A0A0E3TRX4_9CAUD]	101,87698	42,22	1	3	19	0,64102564	2,05405405	0,01317769	0,0422256	30	450	48,644755	5,18603516		
A0A216GV5	Structural protein OS=Salmonella phage BPS1556 OX=2060121 GN=BPS1556_10 PE=4 SV=1 - [A0A216GV5_9CAUD]	86,4972382	88,8	1	9	12	1,92307692	1,29729793	0,14053398	0,09480346	26	125	13,6804712	6,30322266		
Q6KJ00	Peptidase_S49 domain-containing protein OS=Salmonella phage Felix O1 (isolate Felix O1) OX=1283336 PE=3 SV=1 - [Q6KJ00_BPF01]	78,5900934	22,73	21	12	12	2,56410256	1,29729793	0,05699082	0,02882921	25	418	44,9939788	4,76822266		
A0A0D4A12	Uncharacterized protein OS=Salmonella phage vB_SpUM_SP116 OX=1567025 GN=HB2014_72 PE=4 SV=1 - [A0A0D4A12_9CAUD]	74,4387329	41,54	10	12	12	2,56410256	1,29729793	0,06916041	0,03499143	23	325	37,0747177	8,89501953		
A0A029AZK3	Baseplate assembly protein OS=Shigella phage Sf18 OX=2024319 GN=Sf18_gp136 PE=4 SV=1 - [A0A029AZK3_9CAUD]	71,9664409	19,5	2	6	6	0,42735043	0,64864865	0,01230859	0,01868244	22	318	34,7196954	6,77099609		
A0A0D4DAW	Uncharacterized protein OS=Salmonella phage vB_SpUM_SP116 OX=1567025 GN=HB2014_70 PE=4 SV=1 - [A0A0D4DAW4_9CAUD]	63,7099726	33,96	4	1	8	0,21367521	0,86486486	0,00737838	0,02986449	20	265	28,9596408	8,22119141		
A0A0E3XDF0	Uncharacterized protein OS=Escherichia phage vB_EcoM_AYO145A OX=1636202 GN=AYO145A_070 PE=4 SV=1 - [A0A0E3XDF0_9CAUD]	55,2122338	43,36	7	5	5	1,06837607	0,54054054	0,08069548	0,04082755	16	113	13,239602	4,81787109		
A0A386K515	Uncharacterized protein OS=Enterobacter phage phiE3_307 OX=2340711 GN=LINGLNFE_00086 PE=4 SV=1 - [A0A386K515_9CAUD]	54,1603898	7,29	6	1	3	0,21367521	0,32432432	0,00391651	0,00594463	18	480	54,557573	5,82080078		
A0A0F6R8U7	Uncharacterized protein OS=Escherichia coli O157 typing phage 15 OX=1508677 GN=ECP15_0245 PE=4 SV=1 - [A0A0F6R8U7_9CAUD]	54,123055	38,49	4	2	9	0,42735043	0,97297297	0,01493376	0,03400053	18	265	28,6164076	7,23974609		
A0A3T0IN25	Uncharacterized protein OS=Shigella phage vB_SfIM_D04 OX=2500763 GN=vBSfIM004_089 PE=4 SV=1 - [A0A3T0IN25_9CAUD]	51,245755	35,07	8	3	3	0,64102564	0,32432432	0,0435465	0,02203218	13	134	14,7204848	5,17333984		
A0A222Z063	Uncharacterized protein OS=Salmonella phage ST11 OX=2023990 PE=4 SV=1 - [A0A222Z063_9CAUD]	50,5963601	45,27	4	9	9	1,92307692	0,97297297	0,11388288	0,05761858	16	148	16,886442	5,74462891		
S4TPL8	Uncharacterized protein OS=Salmonella phage FSL SP-D12 OX=113771 GN=SP012_00160 PE=4 SV=1 - [S4TPL8_9CAUD]	49,816009	67,86	1	11	11	2,35042735	1,18918919	0,10571867	0,05348793	17	196	22,232851	6,77099609		
A0A5P8D4K9	Uncharacterized protein OS=Pectobacterium phage Wc4-1 OX=2652429 PE=4 SV=1 - [A0A5P8D4K9_9CAUD]	48,404241	10,08	2	1	5	0,21367521	0,54054054	0,00389608	0,00956603	13	486	54,8436643	6,10009766		
A0A0H3UDV1	Uncharacterized protein OS=Escherichia phage HY02 OX=1527531 GN=HY02_072 PE=4 SV=1 - [A0A0H3UDV6_9CAUD]	39,8764544	35,97	6	3	3	0,64102564	0,32432432	0,04104171	0,02076489	11	139	15,6188817	5,94775391		
A0A384TF93	Uncharacterized protein OS=Enterobacteria phage XTG1 OX=1651201 PE=4 SV=1 - [A0A384TF93_9CAUD]	33,2126919	42,86	7	6	6	1,28205128	0,64864865	0,08227905	0,01612875	10	133	15,5817469	6,18896484		
A0A0E3TAX3	Uncharacterized protein OS=Escherichia phage vB_EcoM_AYO145A OX=1636202 GN=AYO145A_063 PE=4 SV=1 - [A0A0E3TAX3_9CAUD]	29,7869463	54,43	12	7	7	1,4957265	0,75675676	0,08464243	0,04284249	11	158	17,6711199	5,04638672		
A0A0A0RLD7	Uncharacterized protein OS=Escherichia phage vB_EcoM_VpaE1 OX=1555238 GN=VpaE1_040 PE=4 SV=1 - [A0A0A0RLD7_9CAUD]	28,8010304	43,09	3	5	5	1,06837607	0,54054054	0,07443868	0,03766195	9	123	14,3524314	8,88037109		
A0A3845J10	Uncharacterized protein OS=Enterobacteria phage XTG1 OX=1651201 PE=4 SV=1 - [A0A3845J10_9CAUD]	27,009408	51	3	4	4	0,85470085	0,43243243	0,07695452	0,03893483	7	100	11,1065711	4,84326172		
O8Q234	Uncharacterized protein OS=Salmonella phage FelixO1 OX=77775 PE=4 SV=1 - [O8Q234_9CAUD]	18,0206006	44,95	19	4	4	0,85470085	0,43243243	0,07096138	0,03590262	5	109	12,0445926	10,081543		
A0A0A0RKH1	Uncharacterized protein OS=Escherichia phage vB_EcoM_VpaE1 OX=1555238 GN=VpaE1_053 PE=4 SV=1 - [A0A0A0RKH1_9CAUD]	14,701772	24,55	9	2	2	0,42735043	0,21621622	0,03660725	0,01852129	5	110	11,6739276	4,70361328		
A0A689WV3	H3JHA OS=Escherichia phage orkinos OX=2696437 GN=orkinos_11 PE=4 SV=1 - [A0A689WV30_9CAUD]	11,4321929	25,44	5	3	3	0,64102564	0,32432432	0,03573192	0,01807842	4	169	17,9398616	5,33837891		
A0A0F6TJH5	Gp138_N domain-containing protein OS=Escherichia coli O157 typing phage 8 OX=1508683 GN=ECP8_01530 PE=4 SV=1 - [A0A0F6TJH5_9CAUD]	8,45222759	12,08	4	2	2	0,42735043	0,21621622	0,01847354	0,00934661	4	207	23,1331056	5,04638672		
A0A689WRQ7	Major capsid protein OS=Escherichia phage ityhuna OX=2696410 GN=ityhuna_34 PE=4 SV=1 - [A0A689WRQ2_9CAUD]	5,95321465	10,25	15	2	2	0,42735043	0,21621622	0,0119398	0,00600849	2	322	35,7920919	5,73193359		
A0A0A0RPQ7	Uncharacterized protein OS=Escherichia phage vB_EcoM_VpaE1 OX=1555238 GN=VpaE1_047 PE=4 SV=1 - [A0A0A0RPQ7_9CAUD]	5,64408278	15,16	52	3	3	0,64102564	0,32432432	0,02506758	0,01268284	3	244	25,571904	4,61474609		
A0A24HYE72	Baseplate J-like protein OS=Salmonella virus VSe102 OX=2053700 GN=vse102_068 PE=4 SV=1 - [A0A24HYE72_9CAUD]	312,923231	66,26	2	23	23	0,42735043	2,48648649	0,00804877	0,04683008	95	489	53,0951105	4,61474609		
Q8XE41	Outer membrane porin C OS=Escherichia coli O157:H7 OX=83334 GN=ompC PE=3 SV=1 - [Q8XE41_9CAUD]	16,782														