

Table 27-1. Virion structure and assembly genes of bacteriophage λ.

| gene | size of protein (amino acids) | in virion? | copies | function, special features |
|------------|-------------------------------|------------|--------|--|
| <i>Nu1</i> | 181 | N | | small terminase subunit: DNA packaging |
| <i>A</i> | 641 | N | | large terminase subunit: DNA packaging |
| <i>W</i> | 68 | Y | ~6 | adaptor between portal and gpFII |
| <i>B</i> | 533 | Y* | 12 | portal (*21 aa cleaved from N-terminus of most subunits) |
| <i>C</i> | 439 | Y* | ~10 | protease (*processed into X1 and X2) |
| <i>Nu3</i> | 131 | N | | scaffolding protein |
| <i>D</i> | 110 | Y | 405 | major capsid decoration protein |
| <i>E</i> | 341 | Y | 405 | major capsid subunit |
| <i>F1</i> | 132 | N | | accessory role in DNA packaging |
| <i>FII</i> | 117 | Y | ~6 | forms tail attachment site on head |
| <i>Z</i> | 192 | ? | | head-tail assembly |
| <i>U</i> | 131 | Y | ~6 | tail shaft stabilization |
| <i>V</i> | 246 | Y | 192 | major tail subunit |
| <i>G</i> | 140 | N | | tail assembly chaperone |
| <i>T</i> | 279* | N | | extension, by translational frameshift, of gpG tail assembly chaperone (*size given is for G-T frameshift product) |
| <i>H</i> | 853* | Y | ~6 | tail length tape measure protein (*~100 aa removed during tail maturation) |
| <i>M</i> | 109 | Y? | | tail tip assembly |
| <i>L</i> | 232 | Y? | | tail tip assembly |
| <i>K</i> | 199 | N? | | tail tip assembly |
| <i>I</i> | 223 | N? | | tail tip assembly |
| <i>J</i> | 1132 | Y | ~3 | tail tip assembly, central tail fiber |
| <i>stf</i> | 774 | Y | 12 | side tail fiber, main structural component |
| <i>tfa</i> | 194 | Y | 12 | side tail fiber, assembly factor and structural component |