

Phage	Transcytosis percent	Capsid size (nm)	Tail size (nm)	Family	Bacterial host	Reference
T3	0.028%	60	23	Podoviridae	<i>Escherichia coli</i>	Fang <i>et al.</i> , <sup>1</sup>
T5	0.0157%	90	250	Siphoviridae	<i>Escherichia coli</i>	Effantin <i>et al.</i> , <sup>2</sup>
T7	0.0013%	60	28	Podoviridae	<i>Escherichia coli</i>	Cuervo <i>et al.</i> , <sup>3</sup>
SPO1	0.005%	~100	~160	Myoviridae	<i>Bacillus subtilis</i>	Parker &
SPP1	0.0001%	60	191	Siphoviridae	<i>Bacillus subtilis</i>	Eiserling <sup>4</sup>
P22	0.0061%	60	20	Podoviridae	<i>Salmonella typhimurium</i>	Lhuillier <i>et al.</i> , <sup>5</sup> Tang <i>et al.</i> , <sup>6</sup>

## REFERENCES

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