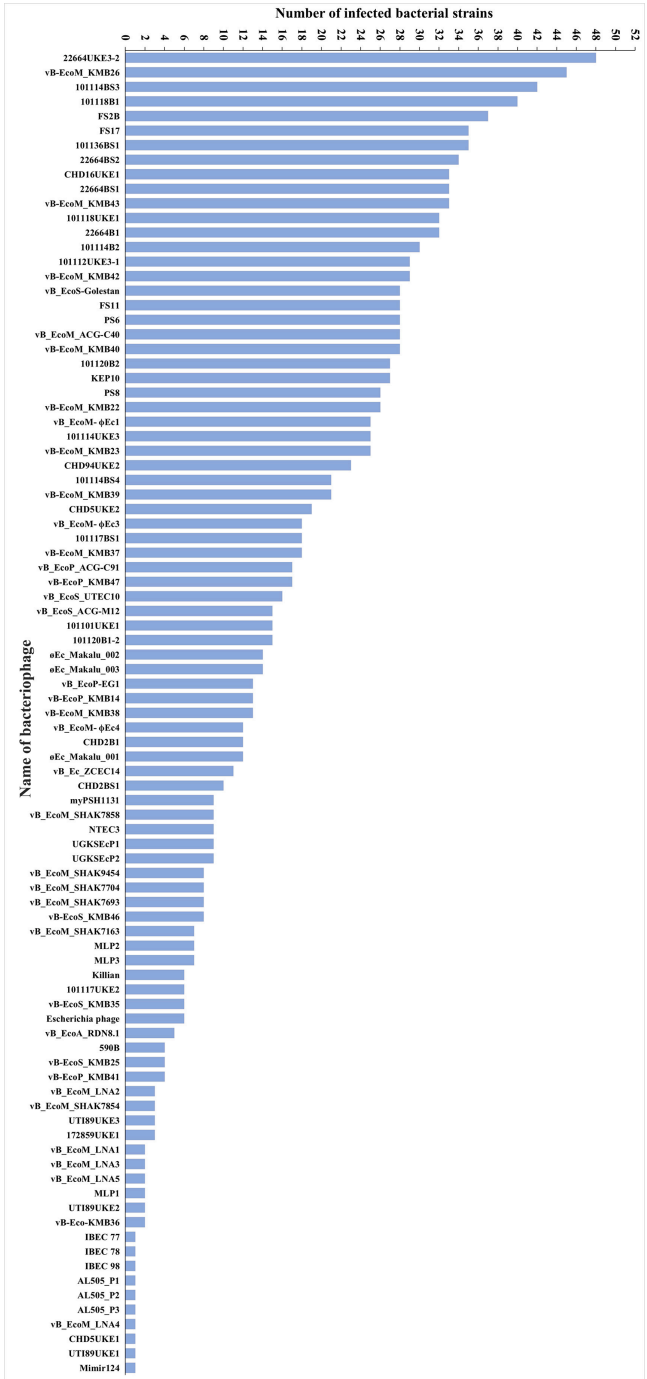

Number of infected bacterial strains
0
2
4
6
8
10
12
14
16
18
20
22
24
26
28
30
32
34
36
38
40
42
44
46
48
50
52
Name of bacteriophage
22664UKE3-2
vB-EcoM_KMB26
101114BS3
101118B1
FS2B
FS17
101136BS1
22664BS2
CHD16UKE1
22664BS1
vB-EcoM_KMB43
101118UKE1
22664B1
101114B2
101112UKE3-1
vB-EcoM_KMB42
vB_EcoS-Golestan
FS11
PS6
vB_EcoM_ACG-C40
vB-EcoM_KMB40
101120B2
KEP10
PS8
vB-EcoM_KMB22
vB_EcoM- ϕEc1
101114UKE3
vB-EcoM_KMB23
CHD94UKE2
101114BS4
vB-EcoM_KMB39
CHD5UKE2
vB_EcoM- ϕEc3
101117BS1
vB-EcoM_KMB37
vB_EcoP_ACG-C91
vB-EcoP_KMB47
vB_EcoS_UTEC10
vB_EcoS_ACG-M12
101101UKE1
101120B1-2
øEc_Makalu_002
øEc_Makalu_003
vB_EcoP-EG1
vB-EcoP_KMB14
vB-EcoM_KMB38
vB_EcoM- ϕEc4
CHD2B1
øEc_Makalu_001
vB_Ec_ZCEC14
CHD2BS1
myPSH1131
vB_EcoM_SHAK7858
NTEC3
UGKSEcP1
UGKSEcP2
vB_EcoM_SHAK9454
vB_EcoM_SHAK7704
vB_EcoM_SHAK7693
vB-EcoS_KMB46
vB_EcoM_SHAK7163
MLP2
MLP3
Killian
101117UKE2
vB-EcoS_KMB35
Escherichia phage
vB_EcoA_RDN8.1
590B
vB-EcoS_KMB25
vB-EcoP_KMB41
vB_EcoM_LNA2
vB_EcoM_SHAK7854
UTI89UKE3
172859UKE1
vB_EcoM_LNA1
vB_EcoM_LNA3
vB_EcoM_LNA5
MLP1
UTI89UKE2
vB-Eco-KMB36
IBEC 77
IBEC 78
IBEC 98
AL505_P1
AL505_P2
AL505_P3
vB_EcoM_LNA4
CHD5UKE1
UTI89UKE1
Mimir124